SEQUENCE LISTING

<110> ROBERTS, JOSEPH SETHURAMAN, NATARAJAN MACALLISTER, THOMAS < 10 > CLONING, OVEREXPRESSION AND THERAPEUTIC USE OF BIOACTIVE HISTIDINE AMMONIA LYASE <130> 078728/0106 <140> 09/833,745 <141> 2001-04-13 <150> 60/197,770 <151> 2000-04-14 <160> 66 <170> PatentIn Ver. 2.1 <210> 1 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic polypeptide <400> 1 Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu Gly 15 Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu Met 25 Gly Glu Gly Glu Ala 35 <210> 2 <211> 40 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic polypeptide <400> 2 Gly Met Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val

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Leu Ala Ser Ala Asp Thr Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly 50 55 60

Ala Leu Ala Thr Arg His Ile Ala Pro Glu Asp Arg Ala Lys Leu Gln 65 70 75 80

Arg Ser Leu Ile Arg Ser His Ala Ala Gly Met Gly Glu Pro Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Ala Lys Thr Leu Ala

Ser Gly Arg Thr Gly Val Arg Pro Val Val Leu Glu Thr Met Val Gly 115 120 125

Met Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu 130 135 140

Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu 145 150 155 160

Met Gly Glu Gly Glu Ala Thr Asp Ala His Gly Asp Ile Arg Pro Val 165 170 175

Pro Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu 180 185 190

Lys Glu Gly Leu Ala Leu Val Asn Gly Thr Asp Gly Met Leu Gly Gln
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Leu Ile Met Ala Leu Ala Asp Leu Asp Glu Leu Leu Asp Ile Ala Asp 210 215 220

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Leu Ala Ser Ala Asp Thr Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly 50 55 60

Ala Leu Ala Thr Arg His Ile Ala Pro Glu Asp Arg Ala Lys Leu Gln 65 70 75 80

Arg Ser Leu Ile Arg Ser His Ala Ala Gly Met Gly Glu Pro Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Ala Lys Thr Leu Ala 100 105 110

Ser Gly Arg Thr Gly Val Arg Pro Val Val Leu Glu Thr Met Val Gly 115 120 125

Met Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu 130 135 140

Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu 145 150 155 160

Met Gly Glu Gly Glu Ala Thr Asp Ala His Gly Asp Ile Arg Pro Val 165 170 175

Pro Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu 180 185 190

Lys Glu Gly Leu Ala Leu Val Asn Gly Thr Asp Gly Met Leu Gly Gln 200 Leu Ile Met Ala Leu Ala Asp Leu Asp Glu Leu Leu Asp Ile Ala Asp Ala Thr Ala Ala Met Ser Val Glu Ala Gln Leu Gly Thr Asp Gln Val Phe Arg Ala Glu Leu His Glu Pro Leu Arg Pro His Pro Gly Gln Gly 245 Arg Ser Ala Gln Asn Met Phe Ala Phe Leu Ala Asp Ser Pro Ile Val 265 Ala Ser His Arg Glu Gly Asp Gly Arg Val Gln Asp Ala Tyr Ser Leu 275 280 285 Arg Cys Ser Pro Gln Val Thr Gly Ala Ala Arg Asp Thr Ile Ala His 295 Ala Arg Leu Val Ala Thr Arg Glu Leu Ala Ala Ile Asp Asn Pro 310 315 Val Val Leu Pro Ser Gly Glu Val Thr Ser Asn Gly Asn Phe His Gly Ala Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Ala Val Ala Asp Leu Gly Ser Ile Ala Glu Arg Arg Thr Asp Arg Met Leu Asp Pro Ala 360 Arg Ser Arg Asp Leu Pro Ala Phe Leu Ala Asp Asp Pro Gly Val Asp 375 Ser Gly Met Met Ile Ala Gln Tyr Thr Gln Ala Gly Leu Val Ala Glu 395 Asn Lys Arg Leu Ala <210> 5 <211> 513 <212> PRT <213> Artificial Sequence

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Leu Ala Ser Ala Asp Thr Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly 50 55 60

Ala Leu Ala Thr Arg His Ile Ala Pro Glu Asp Arg Ala Lys Leu Gln 65 70 75 80

Arg Ser Leu Ile Arg Ser His Ala Ala Gly Met Gly Glu Pro Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Ala Lys Thr Leu Ala 100 105 110

Ser Gly Arg Thr Gly Val Arg Pro Val Val Leu Glu Thr Met Val Gly 115 120 125

Met Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu 130 135 140

Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu 145 150 155 160

Met Gly Glu Gly Glu Ala Thr Asp Ala His Gly Asp Ile Arg Pro Val 165 170 175

Pro Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu 180 185 190

Lys Glu Gly Leu Ala Leu Val Asn Gly Thr Asp Gly Met Leu Gly Gln 195 200 205

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Ala Thr Ala Ala Met Ser Val Glu Ala Gln Leu Gly Thr Asp Gln Val 225 230 235 240

Phe Arg Ala Glu Leu His Glu Pro Leu Arg Pro His Pro Gly Gln Gly 245 250 255

Arg Ser Ala Gln Asn Met Phe Ala Phe Leu Ala Asp Ser Pro Ile Val 260 265 270

Ala Ser His Arg Glu Gly Asp Gly Arg Val Gln Asp Ala Tyr Ser Leu 275 280 285

Arg Cys Ser Pro Gln Val Thr Gly Ala Ala Arg Asp Thr Ile Ala His 290 295 300

Ala Arg Leu Val Ala Thr Arg Glu Leu Ala Ala Ile Asp Asn Pro 305 310 315 320

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Arg Ser Arg Asp Leu Pro Ala Phe Leu Ala Asp Asp Pro Gly Val Asp 370 375 380

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Asn Lys Arg Leu Ala Val Pro Ala Ser Val Asp Ser Ile Pro Ser Ser 405 410 415

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Leu Arg Thr Ser Val Ala Asn Leu Arg Arg Ile Leu Ala Val Glu Met 435 440 445

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Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu Met 145 150 155 160

Gly Glu Gly Glu Ala Thr Asp Ala His Gly Asp Ile Arg Pro Val Pro 165 170 175

Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu Lys 180 185 190

Glu Gly Leu Ala Leu Val Asn Gly Thr Asp Gly Met Leu Gly Gln Leu 195 200 205

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Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu Met 145 150 155 160

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Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu Lys 180 185 190

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- Ser His Arg Glu Gly Asp Gly Arg Val Gln Asp Ala Tyr Ser Leu Arg 275 280 285
- Cys Ser Pro Gln Val Thr Gly Ala Ala Arg Asp Thr Ile Ala His Ala 290 295 300
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- Val Leu Pro Ser Gly Glu Val Thr Ser Asn Gly Asn Phe His Gly Ala 325 330 335
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Leu Ala Xaa Xaa Xaa Pro Val Tyr Gly Xaa Ser Thr Gly Phe Gly 50 55 60

Ala Leu Ala Xaa Arg His Ile Xaa Xaa Glu Xaa Arg Ala Xaa Leu Gln 65 70 75 80

Arg Xaa Xaa Xaa Arg Ser His Ala Ala Gly Met Gly Xaa Xaa Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Xaa Lys Thr Xaa Ala 100 105 110

Ser Gly Xaa Xaa Val Arg Pro Xaa Val Xaa Xaa Thr Met Xaa Gly Xaa 115 120 125

Leu Asn Ala Gly Ile Thr Pro Val Val Xaa Glu Tyr Gly Ser Leu Gly 130 135 140

Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu Met 145 150 155 160

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Thr Ala Ala Xaa Ser Xaa Glu Ala Xaa Leu Gly Thr Asp Xaa Val Xaa 225 230 235 240

Xaa Xaa Glu Leu His Xaa Xaa Xaa Arg Pro His Pro Gly Gln Gly Xaa 250 245 Ser Ala Xaa Asn Met Xaa Xaa Xaa Leu Ala Xaa Ser Xaa Xaa Xaa 265 Xaa His Xaa Xaa Xaa Xaa Arg Val Gln Asp Ala Tyr Ser Xaa Arg Cys Xaa Pro Gln Val Xaa Gly Ala Xaa Arg Asp Thr Xaa Xaa His Ala Xaa Leu Val Ala Xaa Arg Glu Leu Ala Xaa Xaa Asp Asn Pro Val 305 Val Leu Pro Xaa Gly Xaa Val Xaa Ser Asn Gly Asn Phe His Gly Ala Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Xaa Xaa Ala Asp Leu 345 340 Gly Ser Ile Xaa Glu Arg Arg Thr Asp Arg Xaa Leu Asp Xaa Xaa Arg 360 Ser Xaa Xaa Leu Pro Xaa Phe Leu Ala Asp Asp Xaa Gly Val Asp Ser 375 Gly Xaa Met Ile Ala Gln Tyr Thr Gln Ala Xaa Leu Val Xaa Glu Xaa 395 385 Lys Arg Leu Ala Val Pro Ala Xaa Asp Ser Ile Pro Ser Ser Ala Met 410 415 Gln Glu Asp His Val Ser Xaa Gly Trp Xaa Ala Ala Arg Lys Leu Xaa 425 Thr Xaa Val Xaa Asn Leu Xaa Arg Ile Xaa Ala Val Glu Xaa Xaa Xaa Ala Xaa Arg Ala Xaa Xaa Leu Arg Ala Xaa Xaa Xaa Xaa Pro Ala 455 Xaa Xaa Ala Val Xaa Xaa Xaa Leu Arg Xaa Xaa Ala Gly Pro Gly 475 470 Gln Asp Arg Phe Leu Xaa Xaa Xaa Leu Xaa Ala Ala Xaa Xaa Xaa Xaa

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Thr Arg Ile Ala Ser His Asp Leu Glu Asn Leu Gln Arg Ser Leu Val 65 70 75 80

Leu Ser His Ala Ala Gly Ile Gly Ala Pro Leu Asp Asp Asp Leu Val 85 90 95

Arg Leu Ile Met Val Leu Lys Ile Asn Ser Leu Ser Arg Gly Phe Ser 100 105 110

Gly Ile Arg Arg Lys Val Ile Asp Ala Leu Ile Ala Leu Val Asn Ala 115 120 125

Glu Val Tyr Pro His Ile Pro Leu Lys Gly Ser Val Gly Ala Ser Gly 130 135 140

Asp Leu Ala Pro Leu Ala Thr Met Ser Leu Val Leu Leu Gly Glu Gly 145 150 155 160

Lys Ala Arg Tyr Lys Gly Gln Trp Leu Ser Ala Thr Glu Ala Leu Ala 165 170 175 Val Ala Gly Leu Glu Pro Leu Thr Leu Ala Ala Lys Glu Gly Leu Ala 180 185 190

Leu Leu Asn Gly Thr Gln Ala Ser Thr Ala Tyr Ala Leu Arg Gly Leu 195 200 205

Phe Tyr Ala Glu Asp Leu Tyr Ala Ala Ile Ala Cys Gly Gly Leu 210 215 220

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His Glu Ala Arg Gly Gln Arg Gly Gln Ile Asp Thr Ala Ala Cys Phe 245 250 255

Arg Asp Leu Leu Gly Asp Ser Ser Glu Val Ser Leu Ser His Lys Asn 260 265 270

Cys Asp Lys Val Gln Asp Pro Tyr Ser Leu Arg Cys Gln Pro Gln Val 275 280 285

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Pro Phe Leu Val Glu Asn Gly Gly Val Asn Ser Gly Phe Met Ile Ala 370 375 380

Gln Val Thr Ala Ala Ala Leu Ala Ser Glu Asn Lys Ala Leu Ser His 385 390 395 400

Pro His Ser Val Asp Ser Leu Pro Thr Ser Ala Asn Gln Glu Asp His 405 410 415

Val Ser Met Ala Pro Ala Ala Gly Lys Arg Leu Trp Glu Met Ala Glu 420 425 430

Asn Thr Arg Gly Val Pro Ala Ile Glu Trp Leu Gly Ala Cys Gln Gly 435 440 445

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Gly Val Arg Leu Glu Leu Val Arg Leu Ile Glu Ala Met Leu Asp Lys 115 120 125

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Glu Ala Phe Phe Ala Gly Glu Arg Met Lys Gly Asp Ala Ala Leu Lys 165 170 175

Ala Ala Gly Leu Ser Pro Val Thr Leu Ala Ala Lys Glu Gly Leu Ala 180 185 190

Leu Ile Asn Gly Thr Gln Val Ser Thr Ala Leu Ala Leu Ala Gly Leu 195 200 205

Phe Arg Ala His Arg Ala Gly Gln Ala Ala Leu Ile Thr Gly Ala Leu 210 215 220

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Asp Glu Arg Val Gln Asp Pro Tyr Cys Ile Arg Cys Gln Pro Gln Val 275 280 285

Asp Gly Ala Cys Leu Asp Leu Leu Arg Ser Val Ala Ala Thr Leu Thr 290 295 300

Ile Glu Ala Asn Ala Val Thr Asp Asn Pro Leu Val Leu Ser Asp Asn 305 310 315 320

Ser Val Val Ser Gly Gly Asn Phe His Ala Glu Pro Val Ala Phe Ala 325 330 335

Ala Asp Gln Ile Ala Leu Ala Val Cys Glu Ile Gly Ala Ile Ser Gln 340 345 350

Arg Arg Ile Ala Leu Leu Val Asp Pro Ala Leu Ser Leu Arg Leu Pro 355 360 365

Ala Phe Leu Ala Lys Lys Pro Gly Leu Asn Ser Gly Leu Met Ile Ala 370 375 380

Glu Val Thr Ser Ala Ala Leu Met Ser Glu Asn Lys Gln Leu Ser His 385 390 395 400

Pro Ala Ser Val Asp Ser Thr Pro Thr Ser Ala Asn Gln Glu Asp His
405 410 415

Val Ser Met Ala Cys His Gly Ala Arg Arg Leu Leu Gln Met Thr Glu 420 425 430

Asn Leu Phe Ser Ile Ile Gly Ile Glu Ala Leu Ala Ala Val Gln Gly 435 440 445

Ile Glu Phe Arg Ala Pro Leu Thr Thr Ser Pro Glu Leu Gln Lys Ala 450 455 460

Ala Ala Ala Val Arg Gly Val Ser Ser Ser Ile Glu Glu Asp Arg Tyr 465 470 475 480

Met Ala Asp Asp Leu Lys Ala Ala Gly Asp Leu Val Ala Ser Gly Arg 485 490 495

Leu Ala Ala Val Ser Ala Gly Ile Leu Pro Lys Leu Glu Asn 500 505 510

<210> 36

<211> 657

<212> PRT

<213> Mus sp.

<400> 36 Met Pro Arg Tyr Thr Val His Val Arg Gly Glu Trp Leu Ala Val Pro Cys Gln Asp Gly Lys Leu Thr Val Gly Trp Leu Gly Arg Glu Ala Val Arg Arg Tyr Met Lys Asn Lys Pro Asp Asn Gly Gly Phe Thr Ser Val Asp Glu Val Gln Phe Leu Val His Arg Cys Lys Gly Leu Gly Leu Leu Asp Asn Glu Asp Glu Leu Glu Val Ala Leu Glu Asp Asn Glu Phe Val Glu Val Val Ile Glu Gly Asp Val Met Ser Pro Asp Phe Ile Pro Ser Gln Pro Glu Gly Val Phe Leu Tyr Ser Lys Tyr Arg Glu Pro Glu Lys 100 105 Tyr Ile Ala Leu Asp Gly Asp Ser Leu Ser Thr Glu Asp Leu Val Asn 120 Leu Gly Lys Gly Arg Tyr Lys Ile Lys Leu Thr Ser Ile Ala Glu Lys 130 135 Lys Val Gln Gln Ser Arg Glu Val Ile Asp Ser Ile Ile Lys Glu Arg 150 155 Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly Lys Phe Ala Arg Thr 170 Val Ile Pro Ala Asn Lys Leu Gln Glu Leu Gln Val Asn Leu Val Arg Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser Pro Glu Arg Cys Arg Met Leu Leu Ala Leu Arg Ile Asn Val Leu Ala Lys Gly Tyr Ser Gly 215 Ile Ser Leu Glu Thr Leu Lys Gln Val Ile Glu Ala Phe Asn Ala Ser 235 Cys Leu Ser Tyr Val Pro Glu Lys Gly Thr Val Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Gly Leu Ile Gly Glu Gly Lys Met Trp Ser Pro Lys Ser Gly Trp Ala Asp Ala Lys Tyr Val Leu Glu Ala His Gly Leu Lys Pro Ile Val Leu Lys Pro Lys Glu Gly Leu Ala 295

Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu Gly Cys Glu Ala Leu 315 Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp Ile Val Ala Ala Leu 325 Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala Phe Asp Thr Asp Ile His Ala Val Arg Pro His Arg Gly Gln Ile Glu Val Ala Phe Arg Phe Arg Ser Leu Leu Asp Ser Asp His His Pro Ser Glu Ile Ala Glu Ser His Arg Phe Cys Asp Arg Val Gln Asp Ala Tyr Thr Leu Arg Cys Cys 385 390 395 Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe Val Lys Asp 405 410 Ile Ile Thr Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro Met Val Phe 420 425 Ala Ser Arg Gly Glu Thr Ile Ser Gly Gly Asn Phe His Gly Glu Tyr 440 Pro Ala Lys Ala Leu Asp Tyr Leu Ala Ile Gly Val His Glu Leu Ala Ala Ile Ser Glu Arg Arg Ile Glu Arg Leu Cys Asn Pro Ser Leu Ser Glu Leu Pro Ala Phe Leu Val Ala Glu Gly Gly Leu Asn Ser Gly Phe Met Ile Ala His Cys Thr Ala Ala Ala Leu Val Ser Glu Ser Lys Ala 505 Leu Cys His Pro Ser Ser Val Asp Ser Leu Ser Thr Ser Ala Ala Thr 520 Glu Asp His Val Ser Met Gly Gly Trp Ala Ala Arg Lys Ala Leu Arg 535 Val Val Glu His Val Glu Gln Val Leu Ala Ile Glu Leu Leu Ala Ala 555 Cys Gln Gly Ile Glu Phe Leu Arg Pro Leu Lys Thr Thr Pro Leu Glu Lys Val Tyr Asp Leu Val Arg Ser Val Val Arg Pro Trp Ile Lys 585 Asp Arg Phe Met Ala Pro Asp Ile Glu Ala Ala His Arg Leu Leu Leu 595 600

Asp Gln Lys Val Trp Glu Val Ala Ala Pro Tyr Ile Glu Lys Tyr Arg 610 615 620

Met Glu His Ile Pro Glu Ser Arg Pro Leu Ser Pro Thr Ala Phe Ser 625 630 635 640

Leu Glu Ser Leu Arg Lys Asn Ser Ala Thr Ile Pro Glu Ser Asp Asp 645 650 655

Leu

<210> 37

<211> 657

<212> PRT

<213> Rattus sp.

<400> 37

Met Pro Arg Tyr Thr Val His Val Arg Gly Glu Trp Leu Ala Val Pro 1 5 10 15

Cys Gln Asp Gly Lys Leu Ser Val Gly Trp Leu Gly Arg Glu Ala Val 20 25 30

Arg Arg Tyr Met Lys Asn Lys Pro Asp Asn Gly Gly Phe Thr Ser Val\$35\$ 40 45

Asp Glu Val Arg Phe Leu Val Arg Arg Cys Lys Gly Leu Gly Leu Leu 50 55 60

Asp Asn Glu Asp Leu Leu Glu Val Ala Leu Glu Asp Asn Glu Phe Val 65 70 75 80

Glu Val Val Ile Glu Gly Asp Val Met Ser Pro Asp Phe Ile Pro Ser 85 90 95

Gln Pro Glu Gly Val Phe Leu Tyr Ser Lys Tyr Arg Glu Pro Glu Lys 100 105 110

Tyr Ile Ala Leu Asp Gly Asp Ser Leu Ser Thr Glu Asp Leu Val Asn 115 120 125

Leu Gly Lys Gly His Tyr Lys Ile Lys Leu Thr Ser Ile Ala Glu Lys 130 135 140

Lys Val Gln Gln Ser Arg Glu Val Ile Asp Ser Ile Ile Lys Glu Arg 145 150 155 160

Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly Lys Phe Ala Arg Thr
165 170 175

Val Ile Pro Ala Asn Lys Leu Gln Glu Leu Gln Val Asn Leu Val Arg 180 185 190

Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser Pro Glu Arg Cys Arg 195 200 205 ٠,

Met
210Leu
210Leu
AlaLeu
AlaAla
215Asn
AlaVal
AlaLeu
AlaAla
AlaLys
AlaGlu
AlaLys
AlaVal
AlaLys
AlaVal
AlaIle
AlaAla
AlaPhe
AlaAsn
AlaAla
AlaAla
AlaLeu
Ala<

Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu Gly Cys Glu Ala Val 305 310 315 320

Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp Ile Val Ala Ala Leu 325 330 335

Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala Phe Asp Thr Asp Ile 340 345 350

His Ala Val Arg Pro His Arg Gly Gln Ile Glu Val Ala Phe Arg Phe 355 360 365

Arg Ser Leu Leu Asp Ser Asp His His Pro Ser Glu Ile Ala Glu Ser 370 375 380

His Arg Phe Cys Asp Arg Val Gln Asp Ala Tyr Thr Leu Arg Cys Cys 385 390 395 400

Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe Val Lys Asp 405 410 415

Ile Ile Thr Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro Met Val Phe 420 425 430

Ala Ser Arg Gly Glu Thr Ile Ser Gly Gly Asn Phe His Gly Glu Tyr 435 440 445

Pro Ala Lys Ala Leu Asp Tyr Leu Ala Ile Gly Val His Glu Leu Ala 450 455 460

Ala Ile Ser Glu Arg Ile Glu Arg Leu Cys Asn Pro Ser Leu Ser 465 470 475 480

Glu Leu Pro Ala Phe Leu Val Ala Glu Gly Gly Leu Asn Ser Gly Phe 485 490 495

Met Ile Ala His Cys Thr Ala Ala Ala Leu Val Ser Glu Ser Lys Ala 500 505 510

Leu Cys His Pro Ser Ser Val Asp Ser Leu Ser Thr Ser Ala Ala Thr 515 520 525

Glu Asp His Val Ser Met Gly Gly Trp Ala Ala Arg Lys Ala Leu Arg 530 540

Val Ile Glu His Val Glu Gln Val Leu Ala Ile Glu Leu Leu Ala Ala 545 550 555 560

Cys Gln Gly Ile Glu Phe Leu Arg Pro Leu Lys Thr Thr Pro Leu 565 570 575

Glu Lys Val Tyr Asp Leu Val Arg Ser Val Val Arg Pro Trp Ile Lys 580 585 590

Asp Arg Phe Met Ala Pro Asp Ile Glu Ala Ala His Arg Leu Leu 595 600 605

Asp Gln Lys Val Trp Glu Val Ala Ala Pro Tyr Ile Glu Lys Tyr Arg 610 620

Met Glu His Ile Pro Glu Ser Arg Pro Leu Ser Pro Thr Ala Phe Ser 625 630 635 640

Leu Glu Ser Leu Arg Lys Asn Ser Ala Thr Ile Pro Glu Ser Asp Asp 645 650 655

Leu

<210> 38

<211> 635

<212> PRT

<213> Homo sapiens

<400> 38

Met Pro Arg Tyr Thr Val His Val Arg Gly Glu Trp Leu Ala Val Pro 1 5 10 15

Cys Gln Asp Ala Gln Leu Thr Val Gly Trp Leu Gly Arg Glu Ala Val 20 25 30

Arg Arg Tyr Ile Lys Asn Lys Pro Asp Asn Gly Gly Phe Thr Ser Val 35 40 45

Asp Asp Ala His Phe Leu Val Arg Arg Cys Lys Gly Leu Gly Leu Leu 50 60

Asp Asn Glu Asp Arg Leu Glu Val Ala Leu Glu Asn Asn Glu Phe Val 65 70 75 80

Glu Val Val Ile Glu Gly Asp Ala Met Ser Pro Asp Phe Ile Pro Ser 85 90 95

Gln Pro Glu Gly Val Tyr Leu Tyr Ser Lys Tyr Arg Glu Pro Glu Lys 100 105 110

Tyr Ile Glu Leu Asp Gly Asp Arg Leu Thr Thr Glu Asp Leu Val Asn 120 Leu Gly Lys Gly Arg Tyr Lys Ile Lys Leu Thr Pro Thr Ala Glu Lys Arg Val Gln Lys Ser Arg Glu Val Ile Asp Ser Ile Ile Lys Glu Lys Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly Lys Phe Ala Arg Thr Val Ile Pro Ile Asn Lys Leu Gln Glu Leu Gln Val Asn Leu Val Arg 185 Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser Pro Glu Arg Cys Arg 200 Met Leu Leu Ala Leu Arg Ile Asn Val Leu Ala Lys Gly Tyr Ser Gly 215 220 Ile Ser Leu Glu Thr Leu Lys Gln Val Ile Glu Met Phe Asn Ala Ser 230 235 240 Cys Leu Pro Tyr Val Pro Glu Lys Gly Thr Val Gly Ala Ser Gly Asp 250 Leu Ala Pro Leu Ser His Leu Ala Leu Gly Leu Val Gly Glu Gly Lys Met Trp Ser Pro Lys Ser Gly Trp Ala Asp Ala Lys Tyr Val Leu Glu Ala His Gly Leu Lys Pro Val Ile Leu Lys Pro Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu Gly Cys Glu Ala Val 315 Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp Ile Val Ala Ala Leu Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala Phe Asp Thr Asp Ile His Ala Leu Arg Pro His Arg Gly Gln Ile Glu Val Ala Phe Arg Phe Arg Ser Leu Leu Asp Ser Leu Arg Cys Cys Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe Val Lys Asn Ile Ile Thr Thr Glu Leu

Asn Ser Ala Thr Asp Asn Pro Met Val Phe Ala Asn Arg Gly Glu Thr

410

405

Val Ser Gly Gly Asn Phe His Gly Glu Tyr Pro Ala Lys Ala Leu Asp 420 425 430

Tyr Leu Ala Ile Gly Ile His Glu Leu Ala Ala Ile Ser Glu Arg Arg 435 440 445

Ile Glu Arg Leu Cys Asn Pro Ser Leu Ser Glu Leu Pro Ala Phe Leu 450 455 460

Val Ala Glu Gly Gly Leu Asn Ser Gly Phe Met Ile Ala His Cys Thr 465 470 475 480

Ala Ala Ala Leu Val Ser Glu Asn Lys Ala Leu Cys His Pro Ser Ser 490 495

Val Asp Ser Leu Ser Thr Ser Ala Ala Thr Glu Asp His Val Ser Met 500 505 510

Gly Gly Trp Ala Ala Arg Lys Ala Leu Arg Val Ile Glu His Val Glu 515 520 525

Gln Val Leu Ala Ile Glu Leu Leu Ala Ala Cys Gln Gly Ile Glu Phe 530 535 540

Leu Arg Pro Leu Lys Thr Thr Pro Leu Glu Lys Val Tyr Asp Leu 545 550 555 560

Val Arg Ser Val Val Arg Pro Trp Ile Lys Asp Arg Phe Met Ala Pro 565 570 575

Asp Ile Glu Ala Ala His Arg Leu Leu Glu Glu Lys Val Trp Glu
580 585 590

Val Ala Ala Pro Tyr Ile Glu Lys Tyr Arg Met Glu His Ile Pro Glu 595 600 605

Ser Arg Pro Leu Ser Pro Thr Ala Phe Ser Leu Gln Phe Leu His Lys 610 615 620

Lys Ser Thr Lys Ile Pro Glu Ser Glu Asp Leu 625 630 635

<210> 39

<211> 677

<212> PRT

<213> Caenorhabditis elegans

<400> 39

Met Arg Leu Gln Val Gln Ile Gly Thr Glu Cys Val Val Pro Cys
1 5 10 15

Lys Pro Asp Asp Thr Ile His Ala Val Ala Lys Lys Ser Val Glu Lys
20 25 30

Leu Arg Arg Leu Arg Pro Lys Leu Pro Leu Ala Asp Asp Tyr Phe Glu 35 40 45

Val Arg Arg Thr Val Gly Asn Ser Leu Leu Asp Pro Glu Asp Leu Val Ser Asp Val Leu Lys Asp Ser Asp Phe Ile Ile Val Ala Ala Ser Val Glu Glu Thr Glu Asp Ala Lys Glu Ala Lys Lys Gln Glu Glu Ile Asp Asn Ala Arg Ala Glu Ile Glu Lys Ile Asp Asn Arg Arg Arg Lys Val Ser Phe Ala Asp Ser Leu Ala Pro Met Val Leu Ala Pro Pro Thr Lys 120 Leu Leu Ile Leu Asp Gly Asn Ser Leu Leu Pro Glu Asp Leu Val Arg 130 Cys Glu Lys Gly Glu Cys Ala Ile Gln Leu Ser Met Glu Ser Glu Asp 155 Arg Ile Arg Lys Ala Arg Thr Phe Leu Glu Lys Ile Ala Ser Glu His 170 Arg Ala Val Tyr Gly Val Thr Thr Gly Phe Gly Thr Phe Ser Asn Val 185 Thr Ile Pro Pro Glu Lys Leu Lys Lys Leu Gln Leu Asn Leu Ile Arg 200 205 Ser His Ala Thr Gly Tyr Gly Glu Pro Leu Ala Pro Asn Arg Ala Arg Met Leu Leu Ala Leu Arg Ile Asn Ile Leu Ala Lys Gly His Ser Gly 230 Ile Ser Val Glu Asn Ile Lys Lys Met Ile Ala Ala Phe Asn Ala Phe Cys Val Ser Tyr Val Pro Gln Gln Gly Thr Val Gly Cys Ser Gly Asp Leu Cys Pro Leu Ala His Leu Ala Leu Gly Leu Leu Gly Glu Gly Lys

Lys Asn Asn Leu Glu Pro Leu Glu Leu Gly Pro Lys Glu Gly Leu Ala 305

Leu Ile Asn Gly Thr Gln Met Val Thr Ala Leu Gly Ala Tyr Thr Leu 325

Met Trp Ser Pro Thr Thr Gly Trp Gln Pro Ala Asp Val Val Leu Lys

295

Glu Arg Ala His Asn Ile Ala Arg Gln Ala Asp Val Ile Ala Ala Leu 340 345 350 ٠.

Ser Leu Asp Val Leu Lys Gly Thr Thr Arg Ala Tyr Asp Pro Asp Ile 355 360 365

His Arg Ile Arg Pro His Arg Gly Gln Asn Leu Ser Ala Leu Arg Leu 370 375 380

Arg Ala Leu Leu His Ser Glu Ala Asn Pro Ser Gln Ile Ala Glu Ser 385 390 395 400

His Arg Asn Cys Thr Lys Val Gln Asp Ala Tyr Thr Leu Arg Cys Val
405 410 415

Pro Gln Val His Gly Val Val His Asp Thr Ile Glu Phe Val Arg Glu 420 425 430

Ile Ile Thr Thr Glu Met Asn Ser Ala Thr Asp Asn Pro Leu Val Phe 435 440 445

Ala Asp Arg Glu Glu Ile Ile Ser Gly Gly Asn Phe His Gly Glu Tyr 450 455 460

Pro Ala Lys Ala Leu Asp Phe Leu Ala Ile Ala Val Ala Glu Leu Ala 465 470 475 480

Gln Met Ser Glu Arg Arg Leu Glu Arg Leu Val Asn Lys Glu Leu Ser 485 490 495

Gly Leu Pro Thr Glu Leu Thr Pro Asp Gly Gly Leu Asn Ser Gly Phe 500 505 510

Met Thr Val Gln Leu Cys Ala Ala Ser Leu Val Ser Glu Asn Lys Val 515 520 525

Leu Cys His Pro Ser Ser Val Asp Ser Ile Pro Thr Ser Cys Asn Gln 530 535 540

Glu Asp His Val Ser Met Gly Gly Phe Ala Ala Arg Lys Ala Leu Thr 545 550 555 560

Val Val Glu His Val Glu Ala Val Leu Ala Met Glu Leu Leu Ala Ala 565 570 575

Cys Gln Gly Ile Glu Phe Leu Lys Pro Leu Ile Ser Thr Ala Pro Leu 580 585 590

His Lys Ile Tyr Gln Leu Val Arg Ser Val Ala Pro Pro Leu Asn Glu 595 600 605

Asp Arg Tyr Met Lys Pro Glu Ile Asp Ala Val Leu Glu Met Ile Arg 610 615 620

Glu Asn Arg Ile Trp Glu Ala Val Leu Pro His Leu Glu Thr Leu Glu 625 630 635 640

Ala Met Glu Glu Leu Asp Pro Asp Ala Leu Arg Gln Phe Thr Lys Thr 645 650 655

Pro Thr Gly Ile Val Gln Asp Arg Ser Met Ile Pro Ile Ser Asp Asp 660 665 670

Glu Glu Ser Ile Glu 675

<210> 40

<211> 508

<212> PRT

<213> Bacillus subtilis

<400> 40

Met Val Thr Leu Asp Gly Ser Ser Leu Thr Thr Ala Asp Val Ala Arg

1 10 15

Val Leu Phe Asp Phe Glu Glu Ala Ala Ala Ser Glu Glu Ser Met Glu 20 25 30

Arg Val Lys Lys Ser Arg Ala Ala Val Glu Arg Ile Val Arg Asp Glu 35 40 45

Lys Thr Ile Tyr Gly Ile Asn Thr Gly Phe Gly Lys Phe Ser Asp Val 50 55 60

Leu Ile Gln Lys Glu Asp Ser Ala Ala Leu Gln Leu Asn Leu Ile Leu 65 70 75 80

Ser His Ala Cys Gly Val Gly Asp Pro Phe Pro Glu Cys Val Ser Arg 85 90 95

Ala Met Leu Leu Arg Ala Asn Ala Leu Leu Lys Gly Phe Ser Gly
100 105 110

Val Arg Ala Glu Leu Ile Glu Gln Leu Leu Ala Phe Leu Asn Lys Arg 115 120 125

Val His Pro Val Ile Pro Gln Gln Gly Ser Leu Gly Ala Ser Gly Asp 130 135 140

Leu Ala Pro Leu Ser His Leu Ala Leu Ala Leu Ile Gly Gln Gly Glu 145 150 155 160

Val Phe Phe Glu Gly Glu Arg Met Pro Ala Met Thr Gly Leu Lys Lys 165 170 175

Ala Gly Ile Gln Pro Val Thr Leu Thr Ser Lys Glu Gly Leu Ala Leu 180 185 190

Glu Ala Glu Lys Leu Ala Tyr Gln Thr Glu Arg Ile Ala Ser Leu Thr 210 215 220

Ile Glu Gly Leu Gln Gly Ile Ile Asp Ala Phe Asp Glu Asp Ile His 225 230 235 240

Leu Ala Arg Gly Tyr Gln Glu Gln Ile Asp Val Ala Glu Arg Ile Arg 245 250 255

Phe Tyr Leu Ser Asp Ser Gly Leu Thr Thr Ser Gln Gly Glu Leu Arg
260 265 270

Val Gln Asp Ala Tyr Ser Leu Arg Cys Ile Pro Gln Val His Gly Ala 275 280 285

Thr Trp Gln Thr Leu Gly Tyr Val Lys Glu Lys Leu Glu Ile Glu Met 290 295 300

Asn Ala Ala Thr Asp Asn Pro Leu Ile Phe Asn Asp Gly Asp Lys Val 305 310 315 320

Ile Ser Gly Gly Asn Phe His Gly Gln Pro Ile Ala Phe Ala Met Asp 325 330 335

Phe Leu Lys Ile Ala Ile Ser Glu Leu Ala Asn Ile Ala Glu Arg Arg 340 345 350

Ile Glu Arg Leu Val Asn Pro Gln Leu Asn Asp Leu Pro Pro Phe Leu 355 360 365

Ser Pro His Pro Gly Leu Gln Ser Gly Ala Met Ile Met Gln Tyr Ala 370 375 380

Ala Ala Ser Leu Val Ser Glu Asn Lys Thr Leu Ala His Pro Ala Ser 385 390 395 400

Val Asp Ser Ile Pro Ser Ser Ala Asn Glu Glu Asp His Val Ser Met 405 410 415

Gly Thr Ile Ala Ala Arg His Ala Tyr Gln Val Ile Ala Asn Thr Arg 420 425 430

Arg Val Ile Ala Ile Glu Ala Ile Cys Ala Leu Gln Ala Val Glu Tyr 435 440 445

Arg Gly Ile Glu His Ala Ala Ser Tyr Thr Lys Gln Leu Phe Gln Glu 450 455 460

Met Arg Lys Val Val Pro Ser Ile Gln Gln Asp Arg Val Phe Ser Tyr 465 470 475 480

Asp Ile Glu Arg Leu Thr Asp Trp Leu Lys Lys Glu Ser Leu Ile Pro 485 490 495

Asp His Gln Asn Lys Glu Leu Arg Gly Met Asn Ile 500 505

<210> 41

<211> 516

<212> PRT

<213> Streptomyces griseus

<400> 41
Met Asp Met His Thr Val Val Val Gly Thr Ser Gly Thr Thr Ala Glu
1 5 10 15

Asp Val Val Ala Val Ala Arg His Gly Ala Arg Val Glu Leu Ser Ala 20 25 30

Ala Ala Val Glu Ala Leu Ala Ala Ala Arg Leu Ile Val Asp Ala Leu $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Ala Ala Lys Pro Glu Pro Val Tyr Gly Val Ser Thr Gly Phe Gly Ala
50 55 60

Leu Ala Ser Arg His Ile Gly Thr Glu Leu Arg Ala Gln Leu Gln Arg 65 70 75 80

Asn Ile Val Arg Ser His Ala Ala Gly Met Gly Pro Arg Val Glu Arg 85 90 95

Glu Val Val Arg Ala Leu Met Phe Leu Arg Leu Lys Thr Val Ala Ser 100 105 110

Gly His Thr Gly Val Arg Pro Glu Val Ala Gln Thr Met Ala Asp Val 115 120 125

Leu Asn Ala Gly Ile Thr Pro Val Val His Glu Tyr Gly Ser Leu Gly 130 135 140

Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Thr Leu Met 145 150 155 160

Gly Glu Gly Glu Ala Glu Gly Pro Asp Gly Thr Val Arg Pro Ala Gly 165 170 175

Glu Leu Leu Ala Ala His Gly Ile Ala Pro Val Glu Leu Arg Glu Lys 180 185 190

Glu Gly Leu Ala Leu Leu Asn Gly Thr Asp Gly Met Leu Gly Met Leu 195 200 205

Val Met Ala Leu Ala Asp Leu Arg Asn Leu Tyr Thr Ser Ala Asp Ile 210 215 220

Thr Ala Ala Leu Ser Leu Glu Ala Leu Leu Gly Thr Asp Lys Val Leu 225 230 235 240

Ala Pro Glu Leu His Ala Ile Arg Pro His Pro Gly Gln Gly Val Ser 245 250 255

Ala Asp Asn Met Ser Arg Val Leu Ala Gly Ser Gly Leu Thr Gly His 260 265 270

His Gln Asp Asp Ala Pro Arg Val Gln Asp Ala Tyr Ser Val Arg Cys

Ala Pro Gln Val Asn Gly Ala Gly Arg Asp Thr Leu Asp His Ala Ala 290 295 300 Leu Val Ala Gly Arg Glu Leu Ala Ser Ser Val Asp Asn Pro Val Val 305 310 315 320

Leu Pro Asp Gly Arg Val Glu Ser Asn Gly Asn Phe His Gly Ala Pro 325 330 335

Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Val Ala Ala Asp Leu Gly 340 345 350

Ser Ile Cys Glu Arg Arg Thr Asp Arg Leu Leu Asp Lys Asn Arg Ser 355 360 365

His Gly Leu Pro Pro Phe Leu Ala Asp Asp Ala Gly Val Asp Ser Gly 370 375 380

Leu Met Ile Ala Gln Tyr Thr Gln Ala Ala Leu Val Ser Glu Met Lys 385 390 395 400

Arg Leu Ala Val Pro Ala Ser Ala Asp Ser Ile Pro Ser Ser Ala Met 405 410 415

Gln Glu Asp His Val Ser Met Gly Trp Ser Ala Ala Arg Lys Leu Arg 420 425 430

Thr Ala Val Asp Asn Leu Ala Arg Ile Val Ala Val Glu Leu Tyr Ala 435 440 445

Ala Thr Arg Ala Ile Glu Leu Arg Ala Ala Glu Gly Leu Thr Pro Ala 450 455 460

Pro Ala Ser Glu Ala Val Val Ala Ala Leu Arg Ala Ala Gly Ala Glu 465 470 475 480

Gly Pro Gly Pro Asp Arg Phe Leu Ala Pro Asp Leu Ala Ala Asp 485 490 495

Thr Phe Val Arg Glu Gly Arg Leu Val Ala Ala Val Glu Pro Val Thr 500 505 510

Gly Pro Leu Ala 515

<210> 42

<211> 511

<212> PRT

<213> Corynebacteriaceae sp.

<400> 42

Met Ala Ser Ala Pro Gln Ile Thr Leu Gly Leu Ser Gly Ala Thr Ala 1 5 10 15

Asp Asp Val Ile Ala Val Ala Arg His Glu Ala Arg Ile Ser Ile Ser 20 25 30

Pro Gln Val Leu Glu Glu Leu Ala Ser Val Arg Ala His Ile Asp Ala 35 40 45

Leu Ala Ser Ala Asp Thr Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly
50 60

Ala Leu Ala Thr Arg His Ile Ala Pro Glu Asp Arg Ala Lys Leu Gln 65 70 75 80

Arg Ser Leu Ile Arg Ser His Ala Ala Gly Met Gly Glu Pro Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Ala Lys Thr Leu Ala 100 105 110

Ser Gly Arg Ser Val Arg Pro Val Val Leu Glu Thr Met Val Gly Met 115 120 125

Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu Gly 130 135 140

Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu Met 145 150 155 160

Gly Glu Gly Glu Ala Thr Asp Ala His Gly Asp Ile Arg Pro Val Pro 165 170 175

Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu Lys 180 185 190

Glu Gly Leu Ala Leu Val Asn Gly Thr Asp Gly Met Leu Gly Gln Leu 195 200 205

Ile Met Ala Leu Ala Asp Leu Asp Glu Leu Leu Asp Ile Ala Asp Ala 210 215 220

Thr Ala Ala Met Ser Val Glu Ala Gln Leu Gly Thr Asp Gln Val Phe 225 230 235 240

Arg Ala Glu Leu His Glu Pro Leu Arg Pro His Pro Gly Gln Gly Arg 245 250 255

Ser Ala Gln Asn Met Phe Ala Phe Leu Ala Asp Ser Pro Ile Val Ala 260 265 270

Ser His Arg Glu Gly Asp Gly Arg Val Gln Asp Ala Tyr Ser Leu Arg 275 280 285

Cys Ser Pro Gln Val Thr Gly Ala Ala Arg Asp Thr Ile Ala His Ala 290 295 300

Arg Leu Val Ala Thr Arg Glu Leu Ala Ala Ala Ile Asp Asn Pro Val 305 310 315 320

Val Leu Pro Ser Gly Glu Val Thr Ser Asn Gly Asn Phe His Gly Ala 325 330 335

Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Ala Val Ala Asp Leu 340 345 350

:.

Gly Ser Ile Ala Glu Arg Arg Thr Asp Arg Met Leu Asp Pro Ala Arg 355 360 365

Ser Arg Asp Leu Pro Ala Phe Leu Ala Asp Asp Pro Gly Val Asp Ser 370 375 380

Gly Met Met Ile Ala Gln Tyr Thr Gln Ala Gly Leu Val Ala Glu Asn 385 390 395 400

Lys Arg Leu Ala Val Pro Ala Val Asp Ser Ile Pro Ser Ser Ala Met 405 410 415

Gln Glu Asp His Val Ser Leu Gly Trp His Ala Ala Arg Lys Leu Pro 420 425 430

Thr Ser Val Ala Asn Leu Arg Arg Ile Leu Ala Val Glu Met Leu Ile 435 440 445

Ala Gly Arg Ala Leu Asp Leu Arg Ala Pro Leu Lys Pro Gly Pro Ala 450 455 460

Thr Gly Ala Val Leu Glu Val Leu Arg Ser Lys Val Ala Gly Pro Gly 465 470 475 480

Gln Asp Arg Phe Leu Ser Ala Glu Leu Glu Ala Ala Tyr Asp Leu Leu 485 490 495

Ala Asn Gly Ser Val His Lys Ala Leu Glu Ala His Leu Pro Glu
500 505 510

<210> 43

<211> 513

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: 983831/HAL

<400> 43

Met Ala Ser Ala Pro Gln Ile Thr Leu Gly Leu Ser Gly Ala Thr Ala 1 5 10 15

Asp Asp Val Ile Ala Val Ala Arg His Glu Ala Arg Ile Ser Ile Ser 20 25 30

Pro Gln Val Leu Glu Glu Leu Ala Ser Val Arg Ala His Ile Asp Ala 35 40 45

Leu Ala Ser Ala Asp Thr Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly 50 60

Ala Leu Ala Thr Arg His Ile Ala Pro Glu Asp Arg Ala Lys Leu Gln 65 70 75 80

Arg Ser Leu Ile Arg Ser His Ala Ala Gly Met Gly Glu Pro Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Ala Lys Thr Leu Ala 105 Ser Gly Arg Thr Gly Val Arg Pro Val Val Leu Glu Thr Met Val Gly 120 Met Leu Asn Ala Gly Ile Thr Pro Val Val Arg Glu Tyr Gly Ser Leu 135 Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Val Leu 155 Met Gly Glu Gly Glu Ala Thr Asp Ala His Gly Asp Ile Arg Pro Val 170 Pro Glu Leu Phe Ala Glu Ala Gly Leu Thr Pro Val Glu Leu Ala Glu Lys Glu Gly Leu Ala Leu Val Asn Gly Thr Asp Gly Met Leu Gly Gln 200 Leu Ile Met Ala Leu Ala Asp Leu Asp Glu Leu Leu Asp Ile Ala Asp Ala Thr Ala Ala Met Ser Val Glu Ala Gln Leu Gly Thr Asp Gln Val Phe Arg Ala Glu Leu His Glu Pro Leu Arg Pro His Pro Gly Gln Gly 250 245 Arg Ser Ala Gln Asn Met Phe Ala Phe Leu Ala Asp Ser Pro Ile Val 265 Ala Ser His Arg Glu Gly Asp Gly Arg Val Gln Asp Ala Tyr Ser Leu 280 285 Arg Cys Ser Pro Gln Val Thr Gly Ala Ala Arg Asp Thr Ile Ala His 295 Ala Arg Leu Val Ala Thr Arg Glu Leu Ala Ala Ile Asp Asn Pro 315 310 Val Val Leu Pro Ser Gly Glu Val Thr Ser Asn Gly Asn Phe His Gly 330 Ala Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Ala Val Ala Asp 345 Leu Gly Ser Ile Ala Glu Arg Arg Thr Asp Arg Met Leu Asp Pro Ala 355 360 Arg Ser Arg Asp Leu Pro Ala Phe Leu Ala Asp Asp Pro Gly Val Asp 375

Ser Gly Met Met Ile Ala Gln Tyr Thr Gln Ala Gly Leu Val Ala Glu

390

395

Asn Lys Arg Leu Ala Val Pro Ala Ser Val Asp Ser Ile Pro Ser Ser 405 410 415

Ala Met Gln Glu Asp His Val Ser Leu Gly Trp His Ala Ala Arg Lys
420 425 430

Leu Arg Thr Ser Val Ala Asn Leu Arg Arg Ile Leu Ala Val Glu Met 435 440 445

Leu Ile Ala Gly Arg Ala Leu Asp Leu Arg Ala Pro Leu Lys Pro Gly 450 455 460

Pro Ala Thr Gly Ala Val Leu Glu Val Leu Arg Ser Lys Val Ala Gly 465 470 475 480

Pro Gly Gln Asp Arg Phe Leu Ser Ala Glu Leu Glu Ala Ala Tyr Asp 485 490 495

Leu Leu Ala Asn Gly Ser Val His Lys Ala Leu Glu Ala His Leu Pro 500 505 510

Ala

<210>. 44

<211> 513

<212> PRT

<213> Streptomyces coelicolor

<400> 44

Met Ala Ser Met His Thr Val Val Val Gly Thr Ser Gly Val Thr Ala
1 5 10 15

Ser Asp Val Leu Ala Val Ala Arg Ala Gly Ala Arg Ile Glu Leu Ser 20 25 30

Glu Glu Ala Val Ala Ala Leu Ala Ala Ala Arg Ser Val Val Asp Ala 35 40 45

Leu Ala Ala Lys Pro Asp Pro Val Tyr Gly Val Ser Thr Gly Phe Gly 50 55 60

Ala Leu Ala Thr Arg His Ile Ser Pro Glu Leu Arg Gly Arg Leu Gln 65 70 75 80

Arg Asn Ile Val Arg Ser His Ala Ala Gly Met Gly Pro Arg Val Glu 85 90 . 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Leu Lys Thr Val Cys 100 105 110

Ser Gly Arg Thr Gly Val Arg Pro Glu Val Ala Gln Thr Met Ala Asp 115 120 125

Val Leu Asn Ala Gly Ile Thr Pro Val Val His Glu Tyr Gly Ser Leu 130 140

Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Thr Leu 145 150 155 160

Met Gly Glu Gly Asp Ala Glu Gly Pro Asp Gly Thr Val Arg Pro Ala 165 170 175

Gly Glu Leu Leu Ala Ala His Gly Ile Ala Pro Val Glu Leu Arg Glu 180 185 190

Lys Glu Gly Leu Ala Leu Leu Asn Gly Thr Asp Gly Met Leu Gly Met 195 200 205

Leu Val Met Ala Leu Ala Asp Leu Asp Thr Leu Tyr Lys Ser Ala Asp 210 215 220

Ile Thr Ala Ala Leu Thr Met Glu Ala Leu Leu Gly Thr Asp Arg Val 225 230 235 240

Leu Ala Pro Glu Leu His Ala Pro Ile Arg Pro His Pro Gly Gln Ala 245 250 255

Ala Ser Ala Ala Asn Met Ala Ala Val Leu Lys Gly Ser Gly Leu Thr 260 265 270

Gly His His Gln Asp Asp Ala Pro Arg Val Gln Asp Ala Tyr Ser Val 275 280 285

Arg Cys Ala Pro Gln Val Ala Gly Ala Gly Arg Asp Thr Met Ala His 290 295 300

Ala Gly Leu Val Ala Glu Arg Glu Leu Ala Ala Ala Val Asp Asn Pro 305 310 315 320

Val Val Leu Pro Asp Gly Arg Val Glu Ser Asn Gly Asn Phe His Gly 325 330 335

Ala Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Val Ala Val Ala Asp 340 345 350

Leu Gly Ser Ile Ala Glu Arg Arg Thr Asp Arg Leu Leu Asp Lys Asn 355 360 365

Arg Ser His Gly Leu Pro Pro Phe Leu Ala Asp Asp Ala Gly Val Asp 370 375 380

Ser Gly Leu Met Ile Ala Gln Tyr Thr Gln Ala Ala Leu Val Gly Glu 385 390 395 400

Leu Lys Arg Leu Ala Val Pro Ala Ser Ala Asp Ser Ile Pro Ser Ser 405 410 415

Ala Met Gln Glu Asp His Val Ser Met Gly Trp Ser Ala Ala Arg Lys 420 425 430

Leu Arg Thr Ala Val Asp Asn Leu Ala Arg Val Ile Ala Val Glu Leu 435 440 445

Tyr Ala Ala Thr Arg Ala Ile Gln Leu Arg Glu Gly Leu Thr Pro Ala 450 455 460

Pro Ala Ser Gln Ala Val Val Glu Ala Val Arg Ala Ala Val Glu Gly 465 470 475 480

Pro Gly Pro Asp Arg His Leu Ala Pro Asp Leu Ala Ala Ala Asp Ala 485 490 495

Phe Val Arg Ala Gly His Leu Val Ala Ala Ala Glu Ser Val Thr Gly 500 505 510

Pro

<210> 45

<211> 513

<212> PRT

<213> Streptomyces griseus

<400> 45

Met Met Asp Met His Thr Val Val Val Gly Thr Ser Gly Thr Thr Ala 1 5 10 15

Glu Asp Val Val Ala Val Ala Arg His Gly Ala Arg Val Glu Leu Ser 20 25 30

Ala Ala Ala Val Glu Ala Leu Ala Ala Ala Arg Leu Ile Val Asp Ala 35 40 45

Leu Ala Ala Lys Pro Glu Pro Val Tyr Gly Val Ser Thr Gly Phe Gly 50 55 60

Ala Leu Ala Ser Arg His Ile Gly Thr Glu Leu Arg Ala Gln Leu Gln 65 70 75 80

Arg Asn Ile Val Arg Ser His Ala Ala Gly Met Gly Pro Arg Val Glu 85 90 95

Arg Glu Val Val Arg Ala Leu Met Phe Leu Arg Leu Lys Thr Val Ala 100 105 110

Ser Gly His Thr Gly Val Arg Pro Glu Val Ala Gln Thr Met Ala Asp 115 120 125

Val Leu Asn Ala Gly Ile Thr Pro Val Val His Glu Tyr Gly Ser Leu 130 135 140

Gly Cys Ser Gly Asp Leu Ala Pro Leu Ser His Cys Ala Leu Thr Leu 145 150 155 160

Met Gly Glu Gly Glu Ala Glu Gly Pro Asp Gly Thr Val Arg Pro Ala 165 170 175

Gly Glu Leu Leu Ala Ala His Gly Ile Ala Pro Val Glu Leu Arg Glu 180 185 190

- Lys Glu Gly Leu Ala Leu Leu Asn Gly Thr Asp Gly Met Leu Gly Met 195 200 205
- Leu Val Met Ala Leu Ala Asp Leu Arg Asn Leu Tyr Thr Ser Ala Asp 210 215 220
- Ile Thr Ala Ala Leu Ser Leu Glu Ala Leu Leu Gly Thr Asp Lys Val 225 230 235 240
- Leu Ala Pro Glu Leu His Ala Pro Ile Arg Pro His Pro Gly Gln Gly
 245 250 255
- Val Ser Ala Asp Asn Met Ser Arg Val Leu Ala Gly Ser Gly Leu Thr 260 265 270
- Gly His His Gln Asp Asp Ala Pro Arg Val Gln Asp Ala Tyr Ser Val 275 280 285
- Arg Cys Ala Pro Gln Val Asn Gly Ala Gly Arg Asp Thr Leu Asp His 290 295 300
- Ala Ala Leu Val Ala Gly Arg Glu Leu Ala Ser Ser Val Asp Asn Pro 305 310 315 320
- Val Val Leu Pro Asp Gly Arg Val Glu Ser Asn Gly Asn Phe His Gly 325 330 335
- Ala Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Val Ala Ala Asp 340 345 350
- Leu Gly Ser Ile Cys Glu Arg Arg Thr Asp Arg Leu Leu Asp Lys Asn 355 360 365
- Arg Ser His Gly Leu Pro Pro Phe Leu Ala Asp Asp Ala Gly Val Asp 370 375 380
- Ser Gly Leu Met Ile Ala Gln Tyr Thr Gln Ala Ala Leu Val Ser Glu 385 390 395 400
- Met Lys Arg Leu Ala Val Pro Ala Ser Ala Asp Ser Ile Pro Ser Ser 405 410 415
- Ala Met Gl
n Glu Asp His Val Ser Met Gly Tr
p Ser Ala Ala Arg Lys 420 425 430
- Leu Arg Thr Ala Val Asp Asn Leu Ala Arg Ile Val Ala Val Glu Leu 435 440 445
- Tyr Ala Ala Thr Arg Ala Ile Glu Leu Arg Ala Ala Leu Thr Pro Ala 450 455 460
- Pro Ala Ser Glu Ala Val Val Ala Ala Leu Arg Ala Ala Gly Ala Gly 465 470 475 480
- Pro Gly Pro Asp Arg Phe Leu Ala Pro Asp Leu Ala Ala Ala Asp Thr 485 490 495

Phe Val Arg Glu Gly Arg Leu Val Ala Ala Val Glu Pro Val Thr Gly 500 505 510

Pro

<210> 46

<211> 513

<212> PRT

<213> Deinococcus radiodurans

<400> 46

Met Ala Ser Ala Pro Gln Ile Met Ile Leu Asp Arg Asp Leu Asn Leu 1 5 10 15

Glu Gln Phe Ile Ser Val Val Arg His Gly Glu Gln Val Glu Leu Ser 20 25 30

Ala Ala Arg Glu Arg Ile Ala Arg Ala Arg Thr Val Ile Glu Gln
35 40 45

Ile Val Glu Gly Asp Thr Pro Ile Tyr Gly Val Asn Thr Gly Phe Gly 50 60

Lys Phe Glu Asn Val Gln Ile Asp Arg Ser Gln Leu Ala Gln Leu Gln 65 70 75 80

His Asn Leu Ile Val Ser His Ala Ile Gly Met Gly Glu Pro Leu Pro 85 90 95

Ala Glu Val Val Arg Gly Met Leu Leu Leu Arg Ala Gln Ser Leu Ser 100 105 110

Leu Gly His Ser Gly Val Arg Val Glu Val Val Glu Leu Leu Ala 115 120 125

Leu Leu Asn Ala Asp Ala Leu Pro Val Val Pro Ser Gln Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Leu Ala Leu Gly Leu 145 150 155 160

Ile Gly Leu Gly Asp Ile Thr Glu Tyr Gln Gly Gln Val Arg Pro Ala 165 170 175

Ala Asp Val Leu Ala Glu Leu Gly Leu Ser Pro Val Gln Leu Gln Ala 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Leu Met Gly Ser Leu 195 200 205

Leu Ala Leu Ala Leu His Asp Ala Gln Val Leu Leu Gly Thr Ala Asn 210 215 220

Leu Ala Ala Ala Met Thr Val Glu Ala Arg Tyr Gly Ser His Arg Pro 225 230 235 240

Phe Gln Pro Asp Val His Val Gly Leu Arg Pro His Pro Gly Ala Leu 245 250 255

Ala Val Ala Ala Glu Leu Arg Glu Phe Leu Ala Gly Ser Glu Ile Ala 260 265 270

Pro Ser His Leu Thr Gly Asp Gly Lys Val Gln Asp Ala Tyr Ser Leu 275 280 285

Arg Ala Val Pro Gln Val His Gly Ala Thr Trp Asp Ala Leu Ala Gln 290 295 300

Ala Glu Arg Val Leu Ala Val Glu Phe Ala Ser Val Thr Asp Asn Pro 305 310 315 320

Leu Ile Phe Pro Thr Gly Glu Val Val Ser Gly Gly Asn Phe His Gly 325 330 335

Gln Pro Leu Ala Val Thr Ile Asp Ala Leu Lys Val Ala Val Ala Glu 340 345 350

Leu Gly Ser Ile Ser Glu Arg Arg Thr Glu Gln Leu Leu Asn Pro Ala 355 360 365

Leu Ser Arg Gly Leu Pro Ala Phe Leu Thr Pro Asn Gly Gly Leu Asn 370 375 380

Ser Gly Phe Met Ile Ala Gln Tyr Thr Ser Ala Ala Leu Val Ser Glu 385 390 395 400

Asn Lys Val Leu Ser His Pro Ala Ser Val Asp Ser Ile Pro Thr Ser 405 410 415

Ala As
n Glu Asp His Val Ser Met Gly Ala His Ala A
rg Gl
n 420 425 430

Leu Arg Gln Ile Val Ala Asn Val Gln Thr Val Leu Ser Ile Glu Leu 435 440 445

Leu Cys Ala Ala Gln Gly Leu Asp Phe Gln Gln Pro Leu Arg Ala Gly 450 455 460

Arg Gly Val Gln Ala Ala Tyr Glu Tyr Val Arg Thr Phe Val Pro Thr 465 470 475 480

Leu Thr Glu Asp Arg Tyr Phe Arg Pro Asp Leu Leu Arg Leu Arg Gly
485 490 495

Glu Leu Val Ser Gly Arg Val Ala Gln Ala Ala Asp Thr Gln Ala Pro 500 505 510

Ala

<210> 47

<211> 513

<212> PRT

<213> Agrobacterium rhizogenes

<400> 47
Met Ala Ser Ala Pro Gln Ile Thr Leu Gly Leu Ser Gly Val Pro Leu
1 5 10 15

His His Leu Ala Asp Ile Tyr Trp Asn Asn Gly Ser Ala Lys Leu Asp 20 25 30

Pro Ser Phe Asp Ala Ala Val Leu Lys Gly Ala Ala Arg Ile Ala Glu 35 40 45

Ile Ala Ala Gly Asn Ala Pro Val Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Lys Leu Ala Ser Ile Lys Ile Asp Ala Ala Asp Leu Ala Thr Leu Gln 65 70 75 80

Arg Asn Leu Ile Leu Ser His Cys Cys Gly Val Gly Ala Pro Leu Pro 85 90 95

Glu Asn Val Val Arg Leu Ile Met Ala Leu Lys Leu Ile Ser Leu Gly
100 105 110

Arg Gly Ala Ser Gly Val Arg Ile Glu Leu Ile Arg Leu Ile Glu Gly
115 120 125

Met Leu Glu Lys Gly Val Ile Pro Val Ile Pro Glu Lys Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Ala Thr Met 145 150 155 160

Met Gly Glu Gly Glu Ala Phe Asp Tyr Gln Gly Val Gln Met Pro Ser 165 170 175

Lys Asp Ala Leu Ala Lys Ala Gly Leu Ser Pro Val Val Leu Ala Ala 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Thr Ser Thr Ala Leu 195 200 205

Ala Leu Ala Gly Leu Phe Arg Ala His Arg Ala Ala Gln Ser Ala Leu 210 220

Val Thr Gly Ala Leu Ser Thr Asp Ala Ala Met Gly Ser Ser Ala Pro 225 230 235 240

Phe His Pro Asp Ile His Thr Pro Leu Arg Gly His Lys Gly Gln Ile 245 250 255

Asp Ala Gly Ser Ala Leu Arg Asn Leu Leu Gln Gly Ser Glu Ile Arg 260 265 270

Glu Ser His Ile Glu Gly Asp Glu Arg Val Gln Asp Pro Tyr Cys Ile 275 280 285 Arg Cys Gln Pro Gln Val Asp Gly Ala Cys Leu Asp Leu Leu Ala Ser 290 295 300

Val Ala Arg Thr Leu Glu Ile Glu Ala Asn Ala Val Thr Asp Asn Pro 305 310 315 320

Leu Val Leu Ser Asp Asn Ser Val Val Ser Gly Gly Asn Phe His Ala 325 330 335

Glu Pro Val Ala Phe Ala Ala Asp Gln Thr Ala Leu Ala Val Cys Glu 340 345 350

Ile Gly Ala Ile Ala Gln Arg Arg Ile Ala Leu Leu Val Asp Pro Ala 355 360 365

Leu Ser Tyr Gly Leu Pro Ala Phe Leu Ser Lys Lys Pro Gly Leu Asn 370 375 380

Ser Gly Leu Met Ile Ala Glu Val Thr Ser Ala Ala Leu Met Ser Glu 385 390 395 400

Asn Lys Gln Met Ser His Pro Ala Ser Val Asp Ser Thr Pro Thr Ser 405 410 415

Ala Asn Gln Glu Asp His Val Ser Met Ala Cys His Gly Ala Arg Arg
420 425 430

Leu Leu Ala Met Thr Asp Asn Leu Phe Gly Ile Leu Gly Ile Glu Ala 435 440 445

Leu Ala Ala Val Gln Gly Val Glu Leu Arg Gly Pro Leu Lys Thr Ser 450 455 460

Pro Glu Leu Glu Lys Ala Ala Ala Val Leu Arg Ser Ala Val Pro Val 465 470 475 480

Leu Glu Asp Asp Arg Tyr Met Ala Thr Asp Leu Lys Ala Ala Ile Glu 485 490 495

Val Val Ala Ser Gly Ala Leu Val Ser Ala Ile Ser Ser Gly Leu Pro 500 505 510

Val

<210> 48

<211> 513

<212> PRT

<213> Agrobacterium rhizogenes

<400> 48

Met Ala Ser Ala Pro Gln Ile Thr Leu Gly Leu Ser Gly Val Pro Leu 1 5 10 15

His His Leu Ala Asp Ile Tyr Trp Asn Asn Gly Ser Ala Lys Leu Asp 20 25 30

Pro Ser Phe Asp Ala Ala Val Leu Lys Gly Ala Ala Arg Ile Ala Glu 35 40 45

Ile Ala Ala Gly Asn Ala Pro Val Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Lys Leu Ala Ser Ile Lys Ile Asp Ala Ala Asp Leu Ala Thr Leu Gln 65 70 75 80

Arg Asn Leu Ile Leu Ser His Cys Cys Gly Val Gly Ala Pro Leu Pro 85 90 95

Glu Asn Val Val Arg Leu Ile Met Ala Leu Lys Leu Ile Ser Leu Gly
100 105 110

Arg Gly Ala Ser Gly Val Arg Ile Glu Leu Ile Arg Leu Ile Glu Gly 115 120 125

Met Leu Glu Lys Gly Val Ile Pro Val Ile Pro Glu Lys Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Ala Thr Met 145 150 155 160

Met Gly Glu Gly Glu Ala Phe Asp Tyr Gln Gly Val Gln Met Pro Ser 165 170 175

Lys Asp Ala Leu Ala Lys Ala Gly Leu Ser Pro Val Val Leu Ala Ala 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Thr Ser Thr Ala Leu 195 200 205

Ala Leu Ala Gly Leu Phe Arg Ala His Arg Ala Ala Gln Ser Ala Leu 210 215 220

Val Thr Gly Ala Leu Ser Thr Asp Ala Ala Met Gly Ser Ser Ala Pro 225 230 235 240

Phe His Pro Asp Ile His Thr Pro Leu Arg Gly His Lys Gly Gln Ile 245 250 255

Asp Ala Gly Ser Ala Leu Arg Asn Leu Leu Gln Gly Ser Glu Ile Arg 260 265 270

Glu Ser His Ile Glu Gly Asp Glu Arg Val Gln Asp Pro Tyr Cys Ile 275 280 285

Arg Cys Gln Pro Gln Val Asp Gly Ala Cys Leu Asp Leu Leu Ala Ser 290 295 300

Val Ala Arg Thr Leu Glu Ile Glu Ala Asn Ala Val Thr Asp Asn Pro 305 310 315 320

Leu Val Leu Ser Asp Asn Ser Val Val Ser Gly Gly Asn Phe His Ala 325 330 335

Glu Pro Val Ala Phe Ala Ala Asp Gln Thr Ala Leu Ala Val Cys Glu 340 345 350

Ile Gly Ala Ile Ala Gln Arg Arg Ile Ala Leu Leu Val Asp Pro Ala 355 360 365

Leu Ser Tyr Gly Leu Pro Ala Phe Leu Ser Lys Lys Pro Gly Leu Asn 370 375 380

Ser Gly Leu Met Ile Ala Glu Val Thr Ser Ala Ala Leu Met Ser Glu 385 390 395 400

Asn Lys Gln Met Ser His Pro Ala Ser Val Asp Ser Thr Pro Thr Ser 405 410 415

Ala Asn Gln Glu Asp His Val Ser Met Ala Cys His Gly Ala Arg Arg 420 425 430

Leu Leu Ala Met Thr Asp Asn Leu Phe Gly Ile Leu Gly Ile Glu Ala 435 440 445

Leu Ala Ala Val Gln Gly Val Glu Leu Arg Gly Pro Leu Lys Thr Ser 450 455 460

Pro Glu Leu Glu Lys Ala Ala Ala Val Leu Arg Ser Ala Val Pro Val 465 470 475 480

Leu Glu Asp Asp Arg Tyr Met Ala Thr Asp Leu Lys Ala Ala Ile Glu 485 490 / 495

Val Val Ala Ser Gly Ala Leu Val Ser Ala Ile Ser Ser Gly Leu Pro 500 505 510

Val

<210> 49

<211> 513

<212> PRT

<213> Bacillus subtilis

<400> 49

Met Ala Ser Ala Pro Met Val Thr Leu Asp Gly Ser Ser Leu Thr Thr 1 5 10 15

Ala Asp Val Ala Arg Val Leu Phe Asp Phe Glu Glu Ala Ala Ala Ser 20 25 30

Glu Glu Ser Met Glu Arg Val Lys Lys Ser Arg Ala Ala Val Glu Arg

Ile Val Arg Asp Glu Lys Thr Ile Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Lys Phe Ser Asp Val Leu Ile Gln Lys Glu Asp Ser Ala Ala Leu Gln 65 70 75 80

Leu Asn Leu Ile Leu Ser His Ala Cys Gly Val Gly Asp Pro Phe Pro 85 90 95

Glu Cys Val Ser Arg Ala Met Leu Leu Leu Arg Ala Asn Ala Leu Leu 100 105 110

Lys Gly Phe Ser Gly Val Arg Ala Glu Leu Ile Glu Gln Leu Leu Ala 115 120 125

Phe Leu Asn Lys Arg Val His Pro Val Ile Pro Gln Gln Gly Ser Leu 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Ala Leu 145 150 155 160

Ile Gly Gln Gly Glu Val Phe Asp Phe Glu Gly Glu Arg Met Pro Ala 165 170 175

Met Thr Gly Leu Lys Lys Ala Gly Ile Gln Pro Val Thr Leu Thr Ser 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Ala Met Thr Ala Met 195 200 205

Gly Val Val Ala Tyr Ile Glu Ala Glu Lys Leu Ala Tyr Gln Thr Glu 210 215 220

Arg Ile Ala Ser Leu Thr Ile Glu Gly Leu Gln Gly Ile Ile Asp Ala 225 230 235 240

Phe Asp Glu Asp Ile His Leu Ala Leu Arg Gly Tyr Gln Glu Gln Ile 245 250 255

Asp Val Ala Glu Arg Ile Arg Phe Tyr Leu Ser Asp Ser Gly Leu Thr 260 265 270

Thr Ser His Arg Gln Gly Glu Leu Arg Val Gln Asp Ala Tyr Ser Leu 275 280 285

Arg Cys Ile Pro Gln Val His Gly Ala Thr Trp Gln Thr Leu Gly Tyr 290 295 300

Val Lys Glu Lys Leu Glu Ile Glu Met Asn Ala Ala Thr Asp Asn Pro 305 310 315 320

Leu Ile Phe Asn Asp Gly Asp Val Ile Ser Gly Gly Asn Phe His Gly 325 330 335

Gln Pro Ile Ala Phe Ala Met Asp Phe Leu Lys Ile Ala Ile Ser Glu 340 345 350

Leu Ala As
n Ile Ala Glu Arg Arg Ile Glu Arg Leu Val As
n Pro Gl
n 355 360 365

Leu Asn Arg Asp Leu Pro Pro Phe Leu Ser Pro His Pro Gly Leu Gln 370 375 380

Ser Gly Ala Met Ile Met Gln Tyr Ala Ala Ala Ser Leu Val Ser Glu 385 390 395 400

Asn Lys Thr Leu Ala His Pro Ala Ser Val Asp Ser Ile Pro Ser Ser 405 410 415

Ala Asn Gln Glu Asp His Val Ser Met Gly Thr Ile Ala Ala Arg His 420 425 430

Ala Tyr Gln Val Ile Ala Asn Thr Arg Arg Val Ile Ala Ile Glu Ala 435 440 445

Ser Tyr Thr Lys Gln Leu Phe Gln Glu Met Arg Lys Val Val Pro Ser 465 470 475 480

Ile Gln Gln Asp Arg Val Phe Ser Tyr Asp Ile Glu Arg Leu Thr Asp 485 490 495

Trp Leu Lys Lys Glu Ser Leu Ile Pro Asp His Gln Asn Lys Glu Leu 500 505 510

Arg

<210> 50

<211> 513

<212> PRT

<213> Vibrio cholerae

<400> 50

Met Ala Ser Met Leu His Leu Met Ile Lys Pro Gly Gln Leu Ser Leu 1 5 10 15

Lys Gln Leu Arg Gln Val Ser Arg Ser Pro Val Val Leu Ser Leu Asp 20 25 30

Pro Glu Ala Ile Pro Ala Ile Ala Glu Ser Ala Gln Val Val Glu Gln 35 40 45

Val Ile Ser Glu Gly Arg Thr Val Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Leu Leu Ala Asn Thr Lys Ile Ala Pro Gln Asp Leu Glu Thr Leu Gln 65 70 75 80

Lys Ser Ile Val Leu Ser His Ala Ala Gly Ile Gly Glu Leu Met Ser 85 90 95

Asp Glu Thr Val Arg Leu Met Met Leu Leu Lys Ile Asn Ser Leu Ala 100 105 110

Arg Gly Tyr Ser Gly Ile Arg Leu Glu Val Ile Gln Ala Leu Ile Glu 115 120 125 Leu Val Asn Asn Gln Ile Tyr Pro Cys Val Pro Lys Lys Gly Ser Val Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Thr Val Leu Leu Gly Glu Gly Gln Ala Arg Asp Tyr Asn Gly Lys Ile Ile Ser Gly Leu Glu Ala Met Lys Ile Ala Gly Leu Glu Pro Ile Thr Leu Ala Pro 185 Lys Glu Gly Leu Ala Leu Leu Asn Gly Thr Gln Ala Ser Thr Ala Phe Ala Leu Glu Gly Leu Phe Val Ala Glu Asp Leu Phe Ala Ser Ala Thr 210 215 Val Cys Gly Ala Met Ser Val Glu Ala Ala Leu Gly Ser Arg Arg Pro 230 235 Phe Asp Pro Arg Ile His Arg Pro Val Arg Gly His Arg Thr Gln Met Asp Ala Ala Thr Ala Tyr Arg His Leu Leu Val Ser Ser Glu Ile Gly Gln Ser His Ser Asn Cys Glu Gly Lys Val Gln Asp Pro Tyr Ser Leu Arg Cys Gln Pro Gln Val Met Gly Ala Cys Leu Gln Gln Ile Arg Ser 295 Ala Ala Glu Val Leu Glu Val Glu Ala Asn Ser Val Ser Asp Asn Pro 315 Leu Val Phe Ala Asp Gly Asp Ile Ile Ser Gly Gly Asn Phe His Ala Glu Pro Val Ala Met Ala Ala Asp Asn Leu Ala Leu Ala Ile Ala Glu 345 Ile Gly Ser Leu Ser Glu Arg Arg Met Ala Leu Leu Ile Asp Ser Ala Leu Ser Lys Asp Leu Pro Pro Phe Leu Val Asp Asn Gly Gly Val Asn 375 Ser Gly Phe Met Ile Ala Gln Val Thr Ala Ala Ala Leu Ala Ser Glu 385 390 395 Asn Lys Thr Leu Ala His Pro Ala Ser Val Asp Ser Leu Pro Thr Ser 405 410 Ala Asn Gln Glu Asp His Val Ser Met Ala Thr Phe Ala Ala Arg Arg

425

430

420

Leu Arg Asp Met Gly Glu Asn Thr Arg Gly Ile Leu Ala Val Glu Tyr 435 440 445

Leu Ala Ala Ala Gln Gly Leu Asp Phe Arg Ala Pro Leu Lys Ser Ser 450 455 460

Pro Arg Ile Glu Glu Ala Arg Gln Ile Leu Arg Glu Lys Val Pro Phe 465 470 475 480

Tyr Asp Lys Asp Arg Tyr Phe Ala Pro Asp Ile Glu Lys Ala Asn Ala 485 490 495

Leu Leu Ala Gln Leu Ala Val His Asn Arg Leu Met Pro Asp Gln Leu 500 505 510

Leu

<210> 51

<211> 513

<212> PRT

<213> Pseudomonas aeruginosa

<400> 51

Met Ala Ser Ala Met Ser Leu His Leu Lys Pro Gly Gln Leu Thr Leu 1 5 10 15

Ala Asp Leu Arg Gln Ala Tyr Leu Ala Pro Val Arg Leu Ser Leu Asp 20 25 30

Pro Ser Ala Asp Ala Pro Ile Ala Ala Ser Val Ala Cys Val Glu Asn $35 \hspace{1cm} 40 \hspace{1cm} 45$

Ile Ile Ala Glu Gly Arg Thr Ala Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Leu Leu Ala Ser Thr Arg Ile Ser Pro Ala Asp Leu Glu Lys Leu Gln 65 70 75 80

Arg Ser Ile Val Leu Ser His Ala Ala Gly Val Gly Glu Ala Leu Asp 85 90 95

Asp Ala Met Val Arg Leu Val Met Leu Leu Lys Val Asn Ser Leu Ala 100 105 110

Arg Gly Phe Ser Gly Ile Arg Arg Lys Val Ile Asp Ala Leu Ile Ala 115 120 125

Leu Ile Asn Ala Glu Val Tyr Pro His Ile Pro Leu Lys Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Leu Val Leu 145 150 155 160

Ile Gly Glu Ser Arg Ala Arg His Ala Arg Gly Glu Trp Leu Pro Ala 165 170 175 Ala Glu Ala Leu Ala Val Ala Gly Leu Glu Pro Leu Thr Leu Ala Ala 180 185 190

Lys Glu Gly Leu Ala Leu Leu Asn Gly Thr Gln Val Ser Thr Ala Tyr 195 200 205

Ala Leu Arg Gly Leu Phe Glu Ala Glu Asp Leu Phe Ala Ala Thr 210 215 220

Val Cys Gly Gly Leu Ser Val Glu Ala Met Leu Gly Ser Arg Ala Pro 225 230 235 240

Phe Asp Ala Arg Ile His Ala Ala Leu Arg Gly Gln Arg Gly Gln Ile
245 250 255

Asp Val Ala Ala Ala Tyr Arg Asp Leu Leu Ala Ser Ser Glu Val Ala 260 265 270

Arg Ser His Glu Lys Cys Asp Gly Lys Val Gln Asp Pro Tyr Ser Leu 275 280 285

Arg Cys Gln Pro Gln Val Met Gly Ala Cys Leu Thr Gln Met Arg Gln 290 295 300

Ala Ala Glu Val Leu Glu Ile Glu Ala Asn Ala Val Ser Asp Asn Pro 305 310 315 320

Leu Val Phe Ala Ala Gly Asp Val Ile Ser Gly Gly Asn Phe His Ala 325 330 335

Glu Pro Val Ala Met Ala Ala Asp Asn Leu Ala Leu Ala Leu Ala Glu 340 345 350

Ile Gly Ser Leu Ser Glu Arg Arg Ile Ser Leu Met Met Asp Met His 355 360 365

Met Ser Gln Asp Leu Pro Pro Phe Leu Val Ala Asn Gly Gly Val Asn 370 375 380

Ser Gly Phe Met Ile Ala Gln Val Thr Ala Ala Ala Leu Ala Ser Asp 385 390 395 400

Asn Lys Ala Leu Ala His Pro Ala Ser Val Asp Ser Leu Pro Thr Ser 405 410 415

Ala Asn Gln Glu Asp His Val Ser Met Ala Pro Asn Ala Gly Lys Arg
420 425 430

Leu Trp Ala Met Ala Glu Asn Val Arg Gly Ile Leu Ala Val Glu Trp 435 440 445

Leu Gly Ala Cys Gln Gly Leu Asp Phe Arg Glu Gly Leu Lys Ser Ser 450 455 460

Pro Lys Leu Glu Gln Ala Arg Arg Leu Leu Arg Asp Lys Val Pro Tyr 465 470 475 480 Tyr Gln Glu Asp Arg Phe Phe Ala Pro Asp Ile Glu Ala Ala Ser Gln 485 490 495

Leu Leu Ala Ser Gly Cys Leu Asn Ala Leu Leu Pro Ala Arg Leu Leu 500 505 510

Pro

<210> 52

<211> 513

<212> PRT

<213> Bacillus halodurans

<400> 52

Met Ala Met Thr Asn Leu Lys Leu Leu Asp Gly Arg Ser Leu Ser Leu 1 1 5 10

His Asp Leu His Arg Ile Ile Tyr Glu Gly Glu Thr Val Gly Ala Ser 20 25 30

Asp Glu Ser Met Glu Lys Val Lys Gln Ser Arg Lys Ala Val Glu Gln 35 40 45

Ile Val Ala Asp Glu Lys Ile Ile Tyr Gly Ile Thr Thr Gly Phe Gly 50 55 60

Lys Phe Ser Asp Ile Phe Ile Asp Pro Asp Asp Val Glu Asn Leu Gln 65 70 75 80

His Asn Leu Ile Tyr Ser His Ala Cys Gly Val Gly Ser Pro Phe Pro $85 \\ 90 \\ 95$

Glu Thr Val Ser Arg Thr Met Leu Val Leu Arg Ala As
n Ala Leu Leu 100 105 110

Lys Gly Phe Ser Gly Val Arg Pro Leu Val Ile Glu Arg Leu Leu Ala 115 120 125

Leu Val Asn Ala Asn Ile His Pro Val Ile Pro Gln Gln Gly Ser Leu 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Val Leu 145 150 155 160

Leu Gly Glu Gly Glu Val Phe Asp Tyr Lys Gly Thr Lys Thr Lys Ala 165 170 175

Ser Phe Ala Leu Lys Glu Glu Glu Ile Glu Pro Ile Thr Leu Thr Ala 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Ala Met Thr Ala Met 195 200 205

Gly Val Ile Ala Tyr Leu Glu Ala Glu Lys Leu Ala Phe Gln Ser Glu 210 215 220 Ile Ile Ala Ser Leu Thr Met Glu Gly Leu Arg Gly Ile Ile Asp Ala 225 230 235 240

Phe Asp Glu Gln Ile His Phe Ala Leu Arg Gly Tyr Val Glu Gln Val 245 250 255

Asp Val Ala Arg Arg Met Glu Ser Tyr Leu Gln Asp Ser Gln Leu Thr 260 265 270

Thr Ser His Arg Gln Gly Glu Leu Arg Val Gln Asp Ala Tyr Ser Leu 275 280 285

Arg Cys Ile Pro Gln Val His Gly Ala Thr Trp Gln Thr Leu Arg Tyr 290 295 300

Val Lys Glu Lys Leu Glu Ile Glu Met Asn Ala Ala Thr Asp Asn Pro 305 310 315 320

Leu Ile Phe Asp Asn Gly Gln Val Ile Ser Gly Gly Asn Phe His Gly 325 330 335

Gln Gln Ile Ala Leu Ala Met Asp Phe Leu Gly Ile Ala Met Ala Glu 340 345 350

Leu Ala Asn Ile Ser Glu Arg Arg Ile Glu Arg Leu Val Asn Pro Gln 355 360 365

Leu Asn Arg Asp Leu Pro Pro Phe Leu Ser Ala Ala Pro Gly Val Gln 370 375 380

Ser Gly Val Met Ile Leu Gln Tyr Cys Ala Ala Ser Leu Val Ser Glu 385 390 395 400

Asn Lys Thr Leu Ala His Pro Ala Ser Val Asp Ser Ile Pro Ser Ser 405 410 415

Ala Asn Gln Glu Asp His Val Ser Met Gly Thr Ile Gly Ser Arg His 420 425 430

Ile Cys Ala Met Gln Ala Val Asp Ile Arg Gly Arg Glu Lys Met Ala 450 455 460

Ser Phe Thr Lys Lys Ile Leu Glu Lys Gly Arg Glu His Val Pro Tyr 465 470 475 480

Ile Asp Gln Asp Arg Met Phe Ala Lys Asp Ile Glu Arg Ala Ala Lys 485 490 495

Trp Leu Lys Asp Gly Ser Trp Asp Phe Thr Lys Met Arg Glu Lys Glu 500 505 510

Arg

<210> 53

<211> 513

<212> PRT

<213> Pseudomonas putida

<400> 53

Met Ala Ser Ala Thr Glu Leu Thr Leu Lys Pro Gly Thr Leu Thr Leu 1 5 10 15

Ala Gln Leu Arg Ala Ile His Ala Ala Pro Val Arg Leu Gln Leu Asp 20 25 30

Ala Ser Ala Ala Pro Ala Ile Asp Ala Ser Val Ala Cys Val Glu Glu 35 40 45

Ile Ile Ala Glu Asp Arg Thr Ala Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Leu Leu Ala Ser Thr Arg Ile Ala Ser His Asp Leu Glu Asn Leu Gln 65 70 75 80

Arg Ser Leu Val Leu Ser His Ala Ala Gly Ile Gly Ala Pro Leu Asp 85 90 95

Asp Asp Leu Val Arg Leu Ile Met Val Leu Lys Ile Asn Ser Leu Ser 100 105 110

Arg Gly Phe Ser Gly Ile Arg Arg Lys Val Ile Asp Ala Leu Ile Ala 115 120 125

Leu Val Asn Ala Glu Val Tyr Pro His Ile Pro Leu Lys Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Leu Val Leu 145 150 155 160

Leu Gly Glu Gly Lys Ala Arg Asp Tyr Lys Gly Gln Trp Leu Ser Ala 165 170 175

Thr Glu Ala Leu Ala Val Ala Gly Leu Glu Pro Leu Thr Leu Ala Ala 180 185 190

Lys Glu Gly Leu Ala Leu Leu Asn Gly Thr Gln Ala Ser Thr Ala Tyr 195 200 205

Ala Leu Arg Gly Leu Phe Tyr Ala Glu Asp Leu Tyr Ala Ala Ile 210 215 220

Ala Cys Gly Gly Leu Ser Val Glu Ala Val Leu Gly Ser Arg Ser Pro 225 230 235 240

Phe Asp Ala Arg Ile His Glu Ala Leu Arg Gly Gln Arg Gly Gln Ile 245 250 255

Asp Thr Ala Ala Cys Phe Arg Asp Leu Leu Gly Asp Ser Ser Glu Val 260 265 270 Ser Ser His Lys Asn Cys Asp Gly Lys Val Gln Asp Pro Tyr Ser Leu 275 280 285

Arg Cys Gln Pro Gln Val Met Gly Ala Cys Leu Thr Gln Leu Arg Gln 290 295 300

Ala Ala Glu Val Leu Gly Ile Glu Ala Asn Ala Val Ser Asp Asn Pro 305 310 315 320

Leu Val Phe Ala Ala Gly Asp Val Ile Ser Gly Gly Asn Phe His Ala 325 330 335

Glu Pro Val Ala Met Ala Ala Asp Asn Leu Ala Leu Ala Ile Ala Glu 340 345 350

Ile Gly Ser Leu Ser Glu Arg Arg Ile Ser Leu Met Met Asp Lys His 355 360 365

Met Ser Gln Asp Leu Pro Pro Phe Leu Val Glu Asn Gly Gly Val Asn 370 375 380

Ser Gly Phe Met Ile Ala Gln Val Thr Ala Ala Ala Leu Ala Ser Glu 385 390 395 400

Asn Lys Ala Leu Ser His Pro His Ser Val Asp Ser Leu Pro Thr Ser 405 410 415

Ala Asn Glu Asp His Val Ser Met Ala Pro Ala Ala Gly Lys Arg 420 425 430

Leu Trp Glu Met Ala Glu Asn Thr Arg Gly Val Leu Ala Ile Glu Trp 435 440 445

Leu Gly Ala Cys Gl
n Gly Leu Asp Leu Arg Lys Gly Leu Lys Thr Ser 450
 455
 460

Ala Lys Leu Glu Lys Ala Arg Gln Ala Leu Arg Ser Glu Val Ala His 465 470 475 480

Tyr Asp Arg Asp Arg Phe Phe Ala Pro Asp Ile Glu Lys Ala Val Glu
485 490 495

Leu Leu Ala Lys Gly Ser Leu Thr Gly Leu Leu Pro Ala Gly Leu Pro 500 505 510

Ser

<210> 54

<211> 513

<212> PRT

<213> Rhizobium meliloti

<400> 54

Met Ala Ser Ala Pro Gln Ile Thr Leu Arg Pro Gly Ser Val Pro Leu 1 5 10 15

Ser Asp Leu Glu Thr Ile Tyr Trp Thr Gly Ala Pro Ala Arg Leu Asp 20 25 30

Ala Ala Phe Asp Ala Gly Ile Ala Lys Ala Ala Ala Arg Ile Ala Glu 35 40 45

Ile Val Ala Gly Asn Ala Pro Val Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Lys Leu Ala Ser Ile Lys Ile Asp Ser Ser Asp Val Ala Thr Leu Gln 65 70 75 80

Arg Asn Leu Ile Leu Ser His Cys Cys Gly Val Gly Gln Pro Leu Thr 85 90 95

Glu Asp Ile Val Arg Leu Ile Met Ala Leu Lys Leu Ile Ser Leu Gly
100 105 110

Arg Gly Ala Ser Gly Val Arg Leu Glu Leu Val Arg Leu Ile Glu Ala 115 120 125

Met Leu Asp Lys Gly Val Ile Pro Leu Ile Pro Glu Lys Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ala Ala Val Met 145 150 155 160

Met Gly His Gly Glu Ala Phe Phe Ala Gly Glu Arg Met Lys Gly Asp $165 \hspace{1.5cm} 170 \hspace{1.5cm} 175$

Ala Ala Leu Lys Ala Glu Ala Gly Leu Ser Pro Val Thr Leu Ala Ala 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Val Ser Thr Ala Leu 195 200 205

Ala Leu Ala Gly Leu Phe Arg Ala His Arg Ala Gly Gln Ala Ala Leu 210 215 220

Ile Thr Gly Ala Leu Ser Thr Asp Ala Ala Met Gly Ser Ser Ala Pro 225 230 235 240

Phe His Pro Asp Ile Gln His Cys Ala Ala Ile Arg Ala Arg Ser Thr 245 250 255

Arg Ala Ala Asn Leu Arg Gln Leu Leu Thr Gly Ser Pro Ile Arg 260 265 270

Gln Ser His Ile Glu Gly Asp Glu Arg Val Gln Asp Pro Tyr Cys Ile 275 280 285

Arg Cys Gln Pro Gln Val Asp Gly Ala Cys Leu Asp Leu Leu Arg Ser 290 295 300

Val Ala Ala Thr Leu Thr Ile Glu Ala Asn Ala Val Thr Asp Asn Pro 305 310 315 320 Leu Val Leu Ser Asp Asn Ser Val Val Ser Gly Gly Asn Phe His Ala 325 330 335

Glu Pro Val Ala Phe Ala Ala Asp Gln Ile Ala Leu Ala Val Cys Glu 340 345 350

Ile Gly Ala Ile Ser Gln Arg Arg Ile Ala Leu Leu Val Asp Pro Ala 355 360 365

Leu Ser Leu Arg Leu Pro Ala Phe Leu Ala Lys Lys Pro Gly Leu Asn 370 375 380

Ser Gly Leu Met Ile Ala Glu Val Thr Ser Ala Ala Leu Met Ser Glu 385 390 395 400

Asn Lys Gln Leu Ser His Pro Ala Ser Val Asp Ser Thr Pro Thr Ser 405 410 415

Ala Asn Glu Asp His Val Ser Met Ala Cys His Gly Ala Arg Arg 420 425 430

Leu Leu Gln Met Thr Glu Asn Leu Phe Ser Ile Ile Gly Ile Glu Ala 435 440 445

Leu Ala Ala Val Gln Gly Ile Glu Phe Arg Ala Pro Leu Thr Thr Ser 450 455 460

Pro Glu Leu Gln Lys Ala Ala Ala Ala Val Arg Gly Val Ser Ser Ser 465 470 475 480

Ile Glu Glu Asp Arg Tyr Met Ala Asp Asp Leu Lys Ala Ala Gly Asp 485 490 495

Leu Val Ala Ser Gly Arg Leu Ala Ala Ala Val Ser Ala Gly Leu Pro 500 505 510

Lys

<210> 55

<211> 513

<212> PRT

<213> Pseudomonas aeruginosa

<400> 55

Met Ser Asp Leu Pro Ser Val Val Phe Gly Asp Gly Pro Leu Arg Trp 1 5 10 15

Gln Glu Leu Val Ala Val Ala Arg His Gly Ala Arg Leu Glu Leu Ser 20 25 30

Ala Ala Arp Ala Arg Ile Asp Asn Ala Arg Ala Ile Val Cys Arg 35 40 45

Ile Val Ala Asn Gly Glu Arg Ala Tyr Gly Ile Ser Thr Gly Leu Gly 50 55 60

Ala Leu Cys Asp Val Leu Leu Glu Gly Glu Gln Leu Ala Glu Leu Ser
65 70 75 80

Arg Asn Thr Leu Leu Ser His Ala Cys Gly Val Gly Glu Pro Leu Arg 85 90 95

Asp Glu Gln Thr Arg Ala Ile Ile Cys Ala Ala Val Ala Asn Tyr Ser 100 105 110

Gln Gly Lys Ser Gly Leu Asp Arg Ser Leu Val Glu Gly Leu Leu Ala 115 120 125

Leu Leu Asn His Gly Ile Thr Pro Gln Val Pro Ala Gln Gly Ser Val 130 135 140

Gly Tyr Ser Gly Asp Leu Thr His Met Ala His Val Gly Ile Ala Leu 145 150 155 160

Leu Gly Ile Gly Glu Val Ser Asp Tyr Arg Gly Ser Val Val Pro Ala 165 170 175

Ala Ala Ala Leu Ala Ala Glu Gly Leu Ala Thr Val Arg Leu Gly Ala 180 185 190

Lys Asp Gly Leu Cys Leu Val Asn Gly Thr Pro Cys Met Thr Gly Leu 195 200 205

Ala Cys Leu Ala Leu Asp Asp Ala Gln Arg Leu Ala Gln Trp Ala Asp 210 215 220

Val Ile Gly Ala Met Ser Phe Glu Ala Leu Arg Gly Gln Leu Ala Ala 225 230 235 240

Phe Asp Ala Glu Ile His Val Ala Leu Lys Pro His Pro Gly Met Gln 245 250 255

Arg Val Ala Ala Asn Leu Arg Ala Leu Leu Ala Gly Ser Gln Val Leu 260 265 270

Glu Asn Ala Arg Glu Gly Gly Ile Arg Thr Gln Asp Ala Leu Ser Ile 275 280 285

Arg Ser Ile Pro Gln Ile His Gly Ala Cys Arg Asp Gln Leu Ala His 290 295 300

Ala Arg Gln Ile Glu Thr Arg Glu Leu Asn Ser Ala Thr Asp Asn Pro 305 310 315 320

Leu Leu Gly Thr Pro Glu Val Val Ser Gln Ala Asn Pro His Gly 325 330 335

Glu Ser Val Ala Met Ala Ala Asp Leu Leu Ala Ile Ala Val Ala Glu $340 \hspace{1cm} 345 \hspace{1cm} 350$

Leu Gly Gly Val Ala Glu Arg Arg Leu Asp Arg Leu Val Asn Pro Leu 355 360 365

Val Ser Arg Gly Leu Pro Ala Phe Leu Val Gly Lys Pro Gly Val Asn 370 375 380

Ser Gly Met Met Ile Thr Gln Tyr Val Ala Ala Ser Leu Ala Gly Glu 385 390 395 400

Asn Arg Gln Leu Ala Gln Pro Ala Val Val Asp Asn Phe Val Thr Ser 405 410 415.

Ala Leu Gln Glu Asp His Leu Ser Leu Gly Thr Ser Ala Ala Leu Lys
420 425 430

Leu Gly Arg Ala Leu Glu Asn Leu Arg Arg Ile Leu Ala Ile Glu Tyr 435 440 445

Leu Leu Ala Ala Gln Ala Phe Glu Phe Leu Ala Pro Gln Arg Phe Gly 450 455 460

Gln Gly Thr Ala Ala Ala Trp Gly Ile Leu Arg Glu Arg Val Pro Ala 465 470 475 480

Tyr Asp Thr Asp Arg Trp Leu Ala Pro Asp Ile Ala Ser Ala Ala Ala 485 490 495

Ile Leu Gly Glu Arg Lys Ser Leu Ala Arg Leu Ala Ala Ser Ile Gly
500 505 510

Asp

<210> 56 `

<211> 513

<212> PRT

<213> Homo sapiens

<400> 56

Lys Tyr Arg Glu Pro Glu Lys Tyr Ile Glu Leu Asp Gly Leu Thr Thr 1 5 10 15

Glu Asp Leu Val Asn Leu Gly Lys Gly Arg Tyr Lys Ile Lys Leu Thr 20 25 30

Pro Thr Ala Glu Lys Arg Val Gln Lys Ser Arg Glu Val Ile Asp Ser 35 40 45

Ile Ile Lys Glu Lys Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly 50 60

Lys Phe Ala Thr Arg Thr Val Ile Pro Ile Asn Lys Leu Gln Leu Gln 65 70 75 80

Val Asn Leu Val Arg Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser 85 90 95

Pro Glu Arg Cys Arg Met Leu Leu Ala Leu Arg Ile As
n Val Leu Ala 100 105 110

Lys Gly Tyr Ser Gly Ile Ser Leu Glu Thr Leu Lys Gln Val Ile Glu 115 120 125

Met Phe Asn Ala Ser Cys Leu Pro Tyr Val Pro Glu Lys Gly Thr Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Gly Leu 145 150 155 160

Val Gly Glu Gly Lys Met Trp Ser Pro Lys Ser Gly Trp Ala Asp Ala 165 170 175

Lys Tyr Val Leu Glu Ala His Gly Leu Lys Pro Val Ile Leu Lys Pro 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu 195 200 205

Gly Cys Glu Ala Val Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp 210 215 220

Ile Val Ala Ala Leu Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala 225 230 235 240

Phe Asp Thr Asp Ile His Ala Pro Leu Arg Pro His Arg Gly Gln Ile
245 250 255

Glu Val Ala Phe Arg Phe Arg Ser Leu Leu Ser Asp Ser Glu Ile Ala 260 265 270

Glu Ser His Arg Phe Cys Asp Gly Arg Val Gln Asp Ala Tyr Thr Leu 275 280 285

Arg Cys Cys Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe 290 295 300

Val Lys Asn Ile Ile Thr Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro 305 310 315 320

Met Val Phe Ala Asn Gly Glu Thr Val Ser Gly Gly Asn Phe His Gly 325 330 335

Glu Tyr Pro Ala Lys Ala Leu Asp Tyr Leu Ala Ile Gly Ile His Glu 340 345 350

Leu Ala Ala Ile Ser Glu Arg Arg Ile Glu Arg Leu Cys Asn Pro Ser 355 360 365

Leu Ser Arg Glu Leu Pro Ala Phe Leu Val Ala Glu Gly Gly Leu Asn $370 \hspace{1.5cm} 375 \hspace{1.5cm} 380$

Ser Gly Phe Met Ile Ala His Cys Thr Ala Ala Ala Leu Val Ser Glu 385 390 395 400

Asn Lys Ala Leu Cys His Pro Ser Ser Val Asp Ser Leu Ser Thr Ser 405 410 415

Ala Ala Thr Glu Asp His Val Ser Met Gly Gly Trp Ala Ala Arg Lys 420 425 430

Ala Leu Arg Val Ile Glu His Val Glu Gln Val Leu Ala Ile Glu Leu 435 440 445

Leu Ala Ala Cys Gln Gly Ile Glu Phe Leu Arg Pro Leu Lys Thr Thr 450 460

Thr Pro Leu Glu Lys Val Tyr Asp Leu Val Arg Ser Val Val Arg Pro 465 470 475 480

Trp Ile Lys Asp Arg Phe Met Ala Pro Asp Ile Glu Ala Ala His Arg
485 490 495

Leu Leu Leu Glu Gln Lys Val Trp Glu Val Ala Ala Pro Tyr Ile Glu 500 505 510

Lys

<210> 57

<211> 513

<212> PRT

<213> Caenorhabditis elegans

<400> 57

Val Leu Ala Pro Pro Thr Lys Leu Leu Ile Leu Asp Gly Asn Ser Pro 1 $$ 5 $$ 10 $$ 15

Glu Asp Leu Val Arg Cys Glu Lys Gly Glu Cys Ala Ile Gln Leu Ser 20 25 30

Met Glu Ser Glu Asp Arg Ile Arg Lys Ala Arg Thr Phe Leu Glu Lys 35 40 45

Ile Ala Ser Glu His Arg Ala Val Tyr Gly Val Thr Thr Gly Phe Gly 50 55 60

Thr Phe Ser Asn Val Thr Ile Pro Pro Glu Lys Leu Lys Lys Leu Gln 65 70 75 80

Leu Asn Leu Ile Arg Ser His Ala Thr Gly Tyr Gly Glu Pro Leu Ala 85 90 95

Pro Asn Arg Ala Arg Met Leu Leu Ala Leu Arg Ile Asn Ile Leu Ala 100 105 110

Lys Gly His Ser Gly Ile Ser Val Glu Asn Ile Lys Lys Met Ile Ala 115 120 125

Ala Phe Asn Ala Phe Cys Val Ser Tyr Val Pro Gln Gln Gly Thr Val 130 135 140

Gly Cys Ser Gly Asp Leu Cys Pro Leu Ala His Leu Ala Leu Gly Leu 145 150 155 160

450

Leu Gly Glu Gly Lys Met Trp Ser Pro Thr Thr Gly Trp Gln Pro Ala 170 Asp Val Val Leu Lys Lys Asn Asn Leu Glu Pro Leu Glu Leu Gly Pro 185 Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Met Val Thr Ala Leu Gly Ala Tyr Thr Leu Glu Arg Ala His Asn Ile Ala Arg Gln Ala Asp 215 Val Ile Ala Ala Leu Ser Leu Asp Val Leu Lys Gly Thr Thr Arg Ala 230 Tyr Asp Pro Asp Ile His Arg Pro Ile Arg Pro His Arg Gly Gln Asn 250 245 Leu Ser Ala Leu Arg Leu Arg Ala Leu Leu Asn Pro Ser Gln Ile Ala 260 265 Glu Ser His Arg Asn Cys Thr Gly Lys Val Gln Asp Ala Tyr Thr Leu 280 285 Arg Cys Val Pro Gln Val His Gly Val Val His Asp Thr Ile Glu Phe 295 300 Val Arg Glu Ile Ile Thr Thr Glu Met Asn Ser Ala Thr Asp Asn Pro 315 Leu Val Phe Ala Asp Arg Glu Ile Ile Ser Gly Gly Asn Phe His Gly Glu Tyr Pro Ala Lys Ala Leu Asp Phe Leu Ala Ile Ala Val Ala Glu 345 Leu Ala Gln Met Ser Glu Arg Arg Leu Glu Arg Leu Val Asn Lys Glu 360 Leu Ser Arg Gly Leu Pro Thr Phe Leu Thr Pro Asp Gly Gly Leu Asn 375 Ser Gly Phe Met Thr Val Gln Leu Cys Ala Ala Ser Leu Val Ser Glu 395 Asn Lys Val Leu Cys His Pro Ser Ser Val Asp Ser Ile Pro Thr Ser Cys Asn Gln Glu Asp His Val Ser Met Gly Gly Phe Ala Ala Arg Lys 420 Ala Leu Thr Val Val Glu His Val Glu Ala Val Leu Ala Met Glu Leu 440

Leu Ala Ala Cys Gln Gly Ile Glu Phe Leu Lys Pro Leu Ile Ser Thr

455

Ala Pro Leu His Lys Ile Tyr Gln Leu Val Arg Ser Lys Val Ala Pro 465 470 475 480

Pro Asn Glu Asp Arg Tyr Met Lys Pro Glu Ile Asp Ala Val Leu Glu 485 490 495

Met Ile Arg Glu Asn Arg Ile Trp Glu Ala Val Leu Pro His Leu Glu 500 505 510

Thr

<210> 58

<211> 513

<212> PRT

<213> Thermoplasma acidophilum

<400> 58

Met Ala Ser Ala Pro Met Ile Glu Ile Asp Gly Arg Ser Leu Arg Val 1 5 10 15

Glu Asp Val Tyr Ala Val Ala Val Glu Tyr Asp Arg Val Ser Ile Ser 20 25 30

Asp Asp Thr Leu Lys Ala Val Glu Glu Lys His Glu Ala Phe Leu Lys 35 40 45

Leu Ile Asn Ser Gly Lys Thr Val Tyr Gly Val Asn Thr Gly Phe Gly 50 60

Ser Leu Leu Asn Val His Ile Glu Arg Asp Gln Glu Ile Glu Leu Gln 65 70 75 80

Lys Asn Leu Ile Arg Ser His Ser Ser Gly Val Gly Asp Tyr Leu Glu 85 90 95

Asn Arg Tyr Val Arg Ala Ile Met Ala Val Arg Leu Asn Ser Leu Ala 100 105 110

Ala Gly Tyr Ser Ala Val Ser Ala Asp Leu Leu Asn Met Met Val Glu 115 120 125

Met Leu Asn Arg Asp Val Ile Pro Ala Val Pro Lys Tyr Gly Ser Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Ile Gly Leu Ala Met 145 150 155 160

Met Gly Glu Gly Lys Ala Phe Asp Phe Glu Gly Arg Leu Met Asp Ser 165 170 175

Ala Arg Ala Leu Glu Lys Ala Gly Leu Lys Pro Tyr Gln Phe Lys Glu 180 185 190

Lys Glu Gly Val Ala Leu Ile Asn Gly Thr Ser Phe Met Ser Gly Ile 195 200 205 Leu Ser Ile Ala Val Met Asp Ala His Asp Ile Leu Glu Asn Ala Ile 210 215 220

Arg Ser Ala Leu Leu Ser Phe Glu Ala Leu Gly Gly Thr Ser Lys Ala 225 230 235 240

Phe Thr Pro Trp Ile Leu Gly Ala Leu Arg Pro His Leu Gly Gln Val 245 250 255

Ala Ile Gly Asn Arg Phe Arg Glu Tyr Leu Thr Gly Ser Asp Ile Val260 265 270

Ala Ser Lys Arg Ala Asp Ser Val Lys Val Gln Asp Ala Tyr Thr Leu 275 280 285

Arg Cys Ile Pro Gln Val Tyr Gly Ser Val Ala Asp Val Ile Asp Tyr 290 295 300

Val Glu Asn Val Leu Ser Val Glu Ile Asn Ser Ala Thr Asp Asn Pro 305 310 315 320

Leu Val Phe Asn Gly Glu Glu Val Val Ser Gly Gly Asn Phe His Gly 325 330 335

Glu Pro Val Ala Leu Ala Ala Asp Phe Leu Ala Ile Ala Leu Thr Asp 340 345 350

Leu Gly Asn Met Val Glu Arg Arg Ile Ala Arg Leu Val Asp Thr Asn 355 360 365

Leu Ser Arg Gly Leu Pro Pro Phe Leu Thr Pro Asp Ser Gly Leu Asn 370 375 380

Ser Gly Tyr Met Ile Pro Gln Tyr Thr Ala Ala Ala Leu Cys Asn Arg 385 390 395 400

Asn Lys Val Leu Ala Tyr Pro Ser Ser Ala Asp Thr Ile Pro Thr Ser 405 410 415

Ala Asn Glu Asp His Val Ser Met Gly Ala Thr Gly Ser Leu Lys 420 425 430

Leu Leu Glu Ile Ile Asp Asn Val Arg Tyr Ile Ile Ala Ile Glu Tyr 435 440 445

Leu Leu Gly Ser Gln Ala Leu Glu Phe Thr Asp Lys Leu Gly Met Ser 450 455 460

Pro Ser Thr Arg Lys Ile Tyr Glu Lys Ile Arg Glu Lys Val Glu Lys 465 470 475 480

Leu Asp His Asp Arg Pro Pro Ser Phe Asp Ile Glu Thr Ile Arg Lys
485 490 495

Met Met Asp Lys Lys Glu Phe Ile Ser Ala Leu Pro Ala His Leu Pro 500 505 510

Ala

<210> 59

<211> 513

<212> PRT

<213> Mus sp.

<400> 59

Lys Tyr Arg Glu Pro Glu Lys Tyr Ile Ala Leu Asp Gly Asp Ser Thr 1 1 5 10 15

Glu Asp Leu Val Asn Leu Gly Lys Gly Arg Tyr Lys Ile Lys Leu Thr 20 25 30

Ser Ile Ala Glu Lys Lys Val Gln Gln Ser Arg Glu Val Ile Asp Ser 35 40 45

Ile Ile Lys Glu Arg Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly 50 55 60

Lys Phe Ala Thr Arg Thr Val Ile Pro Ala Asn Lys Leu Gln Leu Gln 65 70 75 80

Val Asn Leu Val Arg Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser 85 90 95

Pro Glu Arg Cys Arg Met Leu Leu Ala Leu Arg Ile Asn Val Leu Ala 100 105 110

Lys Gly Tyr Ser Gly Ile Ser Leu Glu Thr Leu Lys Gln Val Ile Glu
115 120 125

Ala Phe Asn Ala Ser Cys Leu Ser Tyr Val Pro Glu Lys Gly Thr Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Gly Leu 145 150 155 160

Ile Gly Glu Gly Lys Met Trp Ser Pro Lys Ser Gly Trp Ala Asp Ala 165 170 175

Lys Tyr Val Leu Glu Ala His Gly Leu Lys Pro Ile Val Leu Lys Pro 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu 195 200 205

Gly Cys Glu Ala Leu Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp 210 215 220

Ile Val Ala Ala Leu Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala 225 230 235 240

Phe Asp Thr Asp Ile His Ala Pro Val Arg Pro His Arg Gly Gln Ile
245 250 255

Glu Val Ala Phe Arg Phe Arg Ser Leu Leu Ser Asp Ser Glu Ile Ala 260 265 270

Glu Ser His Arg Phe Cys Asp Gly Arg Val Gln Asp Ala Tyr Thr Leu 275 280 285

Arg Cys Cys Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe 290 295 300

Val Lys Asp Ile Ile Thr Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro 305 310 315 320

Met Val Phe Ala Ser Gly Glu Thr Ile Ser Gly Gly Asn Phe His Gly 325 330 335

Glu Tyr Pro Ala Lys Ala Leu Asp Tyr Leu Ala Ile Gly Val His Glu 340 345 350

Leu Ala Ala Ile Ser Glu Arg Arg Ile Glu Arg Leu Cys Asn Pro Ser 355 360 365

Leu Ser Arg Glu Leu Pro Ala Phe Leu Val Ala Glu Gly Gly Leu Asn 370 375 380

Ser Gly Phe Met Ile Ala His Cys Thr Ala Ala Ala Leu Val Ser Glu 385 390 395 400

Ser Lys Ala Leu Cys His Pro Ser Ser Val Asp Ser Leu Ser Thr Ser 405 410 415

Ala Ala Thr Glu Asp His Val Ser Met Gly Gly Trp Ala Ala Arg Lys 420 425 430

Ala Leu Arg Val Val Glu His Val Glu Gln Val Leu Ala Ile Glu Leu 435 440 445

Leu Ala Ala Cys Gln Gly Ile Glu Phe Leu Arg Pro Leu Lys Thr Thr 450 455 460

Thr Pro Leu Glu Lys Val Tyr Asp Leu Val Arg Ser Val Val Arg Pro 465 470 475 480

Trp Ile Lys Asp Arg Phe Met Ala Pro Asp Ile Glu Ala Ala His Arg
485 490 495

Leu Leu Leu Asp Gln Lys Val Trp Glu Val Ala Ala Pro Tyr Ile Glu 500 505 510

Lys

<210> 60

<211> 513

<212> PRT

<213> Mus musculus

Glu Asp Leu Val Asn Leu Gly Lys Gly Arg Tyr Lys Ile Lys Leu Thr 20 25 30

Ser Ile Ala Glu Lys Lys Val Gln Gln Ser Arg Glu Val Ile Asp Ser 35 40 45

Ile Ile Lys Glu Arg Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly 50 55 60

Lys Phe Ala Thr Arg Thr Val Ile Pro Ala Asn Lys Leu Gln Leu Gln 65 70 75 80

Val Asn Leu Val Arg Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser 85 90 95

Pro Glu Arg Cys Arg Met Leu Leu Ala Leu Arg Ile As
n Val Leu Ala 100 105 110

Lys Gly Tyr Ser Gly Ile Ser Leu Glu Thr Leu Lys Gln Val Ile Glu 115 120 125

Ala Phe Asn Ala Ser Cys Leu Ser Tyr Val Pro Glu Lys Gly Thr Val 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Gly Leu 145 150 155 160

Ile Gly Glu Gly Lys Met Trp Ser Pro Lys Ser Gly Trp Ala Asp Ala 165 170 175

Lys Tyr Val Leu Glu Ala His Gly Leu Lys Pro Ile Val Leu Lys Pro 180 185 190

Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu 195 200 205

Gly Cys Glu Ala Leu Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp 210 215 220

Ile Val Ala Ala Leu Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala 225 230 235 240

Phe Asp Thr Asp Ile His Ala Pro Val Arg Pro His Arg Gly Gln Ile
245 250 255

Glu Val Ala Phe Arg Phe Arg Ser Leu Leu Ser Asp Ser Glu Ile Ala 260 265 270

Glu Ser His Arg Phe Cys Asp Gly Arg Val Gln Asp Ala Tyr Thr Leu 275 280 285

Arg Cys Cys Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe 290 295 300

Val Lys Asp Ile Ile Thr Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro 305 310 315 320

Met Val Phe Ala Ser Gly Glu Thr Ile Ser Gly Gly Asn Phe His Gly 325 330 335

Glu Tyr Pro Ala Lys Ala Leu Asp Tyr Leu Ala Ile Gly Val His Glu 340 345 350

Leu Ala Ala Ile Ser Glu Arg Arg Ile Glu Arg Leu Cys Asn Pro Ser 355 360 365

Leu Ser Arg Glu Leu Pro Ala Phe Leu Val Ala Glu Gly Gly Leu Asn 370 375 380

Ser Gly Phe Met Ile Ala His Cys Thr Ala Ala Ala Leu Val Ser Glu 385 390 395 400

Ser Lys Ala Leu Cys His Pro Ser Ser Val Asp Ser Leu Ser Thr Ser 405 410 415

Ala Ala Thr Glu Asp His Val Ser Met Gly Gly Trp Ala Ala Arg Lys 420 425 430

Ala Leu Arg Val Val Glu His Val Glu Gln Val Leu Ala Ile Glu Leu 435 440 445

Leu Ala Ala Cys Gln Gly Ile Glu Phe Leu Arg Pro Leu Lys Thr Thr 450 455 460

Thr Pro Leu Glu Lys Val Tyr Asp Leu Val Arg Ser Val Val Arg Pro 465 470 475 480

Trp Ile Lys Asp Arg Phe Met Ala Pro Asp Ile Glu Ala Ala His Arg 485 490 495

Leu Leu Leu Asp Gln Lys Val Trp Glu Val Ala Ala Pro Tyr Ile Glu 500 505 510

Lys

<210> 61

<211> 513

<212> PRT

<213> Rattus sp.

<400> 61

Lys Tyr Arg Glu Pro Glu Lys Tyr Ile Ala Leu Asp Gly Asp Ser Thr
1 5 10 15

Glu Asp Leu Val Asn Leu Gly Lys Gly His Tyr Lys Ile Lys Leu Thr 20 25 30

Ser Ile Ala Glu Lys Lys Val Gln Gln Ser Arg Glu Val Ile Asp Ser 35 40 45

- Ile Ile Lys Glu Arg Thr Val Val Tyr Gly Ile Thr Thr Gly Phe Gly 50 55 60
- Lys Phe Ala Thr Arg Thr Val Ile Pro Ala Asn Lys Leu Gln Leu Gln 65 70 75 80
- Val Asn Leu Val Arg Ser His Ser Ser Gly Val Gly Lys Pro Leu Ser 85 90 95
- Pro Glu Arg Cys Arg Met Leu Leu Ala Leu Arg Ile Asn Val Leu Ala 100 105 110
- Lys Gly Tyr Ser Gly Ile Ser Leu Glu Thr Leu Lys Gln Val Ile Glu
 115 120 125
- Val Phe Asn Ala Ser Cys Leu Ser Tyr Val Pro Glu Lys Gly Thr Val 130 135 140
- Gly Ala Ser Gly Asp Leu Ala Pro Leu Ser His Leu Ala Leu Gly Leu 145 150 155 160
- Ile Gly Glu Gly Lys Met Trp Ser Pro Lys Ser Gly Trp Ala Asp Ala 165 170 175
- Lys Tyr Val Leu Glu Ala His Gly Leu Lys Pro Ile Val Leu Lys Pro 180 185 190
- Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Met Ile Thr Ser Leu 195 200 205
- Gly Cys Glu Ala Val Glu Arg Ala Ser Ala Ile Ala Arg Gln Ala Asp 210 215 220
- Ile Val Ala Ala Leu Thr Leu Glu Val Leu Lys Gly Thr Thr Lys Ala 225 230 235 240
- Phe Asp Thr Asp Ile His Ala Pro Val Arg Pro His Arg Gly Gln Ile 245 250 255
- Glu Val Ala Phe Arg Phe Arg Ser Leu Leu Ser Asp Ser Glu Ile Ala 260 265 270
- Glu Ser His Arg Phe Cys Asp Gly Arg Val Gln Asp Ala Tyr Thr Leu 275 280 285
- Arg Cys Cys Pro Gln Val His Gly Val Val Asn Asp Thr Ile Ala Phe 290 295 300
- Val Lys Asp Ile Ile Thr Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro 305 310 315 320
- Met Val Phe Ala Ser Gly Glu Thr Ile Ser Gly Gly Asn Phe His Gly 325 330 335
- Glu Tyr Pro Ala Lys Ala Leu Asp Tyr Leu Ala Ile Gly Val His Glu 340 345 350

Leu Ala Ala Ile Ser Glu Arg Arg Ile Glu Arg Leu Cys Asn Pro Ser 355 360 365

Leu Ser Arg Glu Leu Pro Ala Phe Leu Val Ala Glu Gly Gly Leu Asn $370 \hspace{1.5cm} 375 \hspace{1.5cm} 380$

Ser Gly Phe Met Ile Ala His Cys Thr Ala Ala Ala Leu Val Ser Glu 385 390 395 400

Ser Lys Ala Leu Cys His Pro Ser Ser Val Asp Ser Leu Ser Thr Ser 405 410 415

Ala Ala Thr Glu Asp His Val Ser Met Gly Gly Trp Ala Ala Arg Lys 420 425 430

Ala Leu Arg Val Ile Glu His Val Glu Gln Val Leu Ala Ile Glu Leu 435 440 445

Leu Ala Ala Cys Gln Gly Ile Glu Phe Leu Arg Pro Leu Lys Thr Thr 450 455 460

Thr Pro Leu Glu Lys Val Tyr Asp Leu Val Arg Ser Val Val Arg Pro 465 470 475 480

Trp Ile Lys Asp Arg Phe Met Ala Pro Asp Ile Glu Ala Ala His Arg 485 490 495

Leu Leu Leu Asp Gln Lys Val Trp Glu Val Ala Ala Pro Tyr Ile Glu
500 505 510

Lys

<210> 62

<211> 513

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Uncultured bacterium pCosAS1

<400> 62

Met Ala Ser Met Asn Ala Leu Thr Leu Thr Pro Gly Thr Leu Thr Leu 1 5 10 15

Ala Gln Leu Arg Gln Val Trp Gln Gln Pro Leu Gln Leu Thr Leu Asp

Glu Ser Ala His Glu Ala Ile Asn Asp Ser Val Ala Cys Val Glu Ala 35 40 45

Ile Val Ala Glu Gly Arg Thr Ala Tyr Gly Ile Asn Thr Gly Phe Gly 50 55 60

Leu Leu Ala Gln Thr Arg Ile Ala Thr His Asp Leu Glu Asn Leu Gln 65 70 75 80

Arg Ser Leu Val Leu Ser His Ala Ala Gly Val Gly Glu Pro Leu Asp Asp Asp Ile Val Arg Leu Met Wet Val Leu Lys Ile Asn Ser Leu Ala 105 Arg Gly Phe Ser Gly Ile Arg Leu Ser Val Ile Gln Ala Leu Ile Ala Leu Val Asn Ala Gly Val Tyr Ser Val Asp Pro Ala Lys Gly Ser Val Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Leu Thr Leu 150 155 145 Leu Gly Glu Gly Lys Ala Arg Asp Tyr Arg Gly Glu Trp Leu Pro Ala 170 Ala Thr Ala Leu Gln Lys Ala Gly Leu Ala Pro Val Thr Leu Ala Ala 185 180 Lys Glu Gly Leu Ala Leu Leu Asn Gly Thr Gln Ala Ser Thr Ala Phe 200 Ala Leu Arg Gly Leu Phe Glu Ala Glu Asp Leu Phe Ala Ser Ala Val 215 Val Cys Gly Ala Leu Thr Thr Glu Ala Val Leu Gly Ser Arg Arg Pro 235 Phe Asp Ala Arg Ile His Glu Pro Val Arg Gly Gln Arg Gly Gln Ile Asp Ala Ala Leu Phe Arg His Val Leu Thr Asp Thr Ser Ala Ile Ala Ser His His Asn Cys Asp Gly Lys Val Gln Asp Pro Tyr Ser Leu Arg Cys Gln Pro Gln Val Met Gly Ala Cys Leu Thr Gln Met Arg Gln 295 Val Ala Glu Val Leu Leu Val Glu Ser Asn Ala Val Ser Asp Asn Pro 310 315 Leu Val Phe Ala Ala Asn Glu Met Val Phe Arg Gly Asn Phe His Ala 325 Glu Pro Val Ala Met Ala Ala Asp Asn Leu Ala Leu Ala Ile Ala Glu 345 Ile Gly Ala Leu Ser Glu Arg Arg Ile Ala Leu Met Met Asp Lys His 355 Met Ser Gln Asp Leu Pro Pro Phe Leu Val Arg Asn Gly Gly Val Asn 375

Ser Gly Phe Met Ile Ala Gln Val Thr Ala Ala Ala Leu Ala Ser Glu 385 390 395 400

Asn Lys Gly Leu Cys His Pro Thr Ser Val Asp Lys Ile Pro Pro Ser 405 410 415

Ala Asn Gln Glu Asp His Val Ser Met Ala Pro Ala Ala Gly Arg Arg 420 425 430

Leu Trp Glu Met Ala Gly Asn Thr Arg Gly Val Leu Ala 'Val Glu Trp 435 440 445

Leu Ala Ala Cys Gln Gly Ala Asp Leu Arg Asp Gly Leu Thr Ser Ser 450 455 460

Pro Leu Leu Glu Gln Ala Arg Gln Ser Cys Gly Glu Gln Val Ala His 465 470 475 480

Tyr Asp Asp Asp Arg Phe Phe Ala Pro Asp Ile Glu Ala Ala Ile Ser 485 490 495

Leu Leu Asn Lys Gly Ser Leu Val Gly Leu Leu Pro Ala Phe Leu Pro 500 505 510

Ala

<210> 63

<211> 513

<212> PRT

<213> Rhizobium meliloti

<400> 63

Met Ala Ser Ala Met Gly Glu Met Ile Ser Leu Asp Gly Pro Leu Thr 1 5 10 15

Trp Arg Glu Ile Ala Ser Ile Ala Glu Gly Ala Ser Leu Asp Leu Ser 20 25 30

Gly Pro Ala Arg Leu Arg Ile Ala Gln Ala Arg Arg Ile Val Asp Ala 35 40 45

Leu Val Glu Arg Gly Ile Arg Gly Tyr Gly Ile Asn Thr Gly Val Gly 50 55 60

Ala Leu Cys Asp Val Ile Ile Ser Arg Glu Asn Gln Gln Ala Leu Ser 65 70 75 80

Arg Asn Ile Ile Leu Ser His Ala Cys Gly Val Gly Asp Pro Leu Gly 85 90 95

Arg Val Glu Ala Arg Ala Val Met Ala Ala Gln Ile Ala Asn Leu Thr 100 105 110

His Gly Tyr Ser Gly Val Arg Val Glu Thr Ala Glu Met Leu Leu Ala 115 120 125

Leu Leu Asn Ala Asp Ile Ile Pro Leu Ile Pro Ser Arg Gly Ser Val 135 Gly Tyr Ser Gly Asp Leu Ala Pro Leu Thr His Ala Ala Leu Val Leu 155 Ile Gly His Gly Ser Ala Met Gln Gly Thr Glu Arg Leu Ser Gly Ala Asp Ala Leu Phe Ala Arg Leu Gly Leu Ala Pro Leu Arg Leu Glu Ala Lys Glu Gly Leu Ser Leu Val Asn Gly Thr Pro Cys Ala Thr Gly Leu Ala Ala Leu Ala Leu Ala Arg Thr Glu Arg Leu Phe Ala Trp Ala Asp 215 210 Ala Ala Ala Met Thr Tyr Glu Ala Asn Leu Gly Ser Gln Ala Asn 230 235 Ala Phe Ala Glu Leu Pro Leu Ala Leu Arg Gln Ser Pro Gly Leu Ser 245 250 Ala Val Gly Glu Gly Leu Arg Asp Trp Leu Ala Asp Ser Pro Met Leu 265 Ala Gly His Arg Thr Ala Gly Thr Arg Thr Gln Asp Pro Leu Ser Leu 280 285 Arg Ala Val Pro Gln Val His Gly Ala Ala Arg Asp Ala Phe Gly Gln Val Ala Glu Ile Val Asp Arg Glu Leu Ala Ser Val Thr Asp Asn Pro 310 Ala Val Ala Gly Ser Pro Glu Val His Ser Gln Ala His Ala Val Gly 330 Ala Ala Leu Gly Leu Ala Met Asp Ser Leu Ala Val Ala Val Ala Glu 345 Val Ala Ala Ile Ser Glu Arg Arg Ile Asp Arg Leu Val Asn Pro Leu Val Ser Arg Gly Leu Pro Ala Phe Leu Ala Gly Asp Ser Gly Val Ser 375 Ser Gly Phe Met Ile Ala Gln Tyr Thr Ala Ala Ala Leu Val Ala Glu 390 395 Asn Arg Arg Leu Ala Ala Pro Ala Ser Leu Asp Gly Gly Ile Thr Ser 410 Ala Leu Gln Glu Asp Met Leu Thr His Ala Thr Pro Ala Ala Trp Lys 425 420

Ala Leu Ser Ile Val Asp Asn Leu Glu Arg Ile Leu Ala Ile Glu Leu 435 440 445

Leu Ala Ala His Arg Pro Met Ser Cys Ser Arg Lys Arg Arg Ala Arg 450 455 460

Arg Asn Ala Pro Leu Pro Phe Thr Gly Thr Tyr Ala Arg Arg Ser Pro 465 470 475 480

Pro Ile Ala Thr Ile Val Arg Ser Ala Glu Leu Glu Ala Ala Tyr Asp 485 490 495

Leu Leu Ala Asn Gly Ser Val His Lys Ala Leu Glu Ala His Leu Pro 500 505 510

Ala

<210> 64

<211> 513

<212> PRT

<213> Halobacterium sp.

<400> 64

Met Ala Ser Ala Pro Gln Ile Thr Leu Gly Leu Ser Gly Ala Thr Ala 1 5 10 15

Asp Asp Val Ile Ala Val Ala Arg His Glu Ala Arg Ile Ser Ile Ser 20 25 30

Pro Gln Val Leu Glu Glu Leu Ala Ser Val Arg Ala His Ile Asp Ala 35 40 45

Leu Ala Ser Ala Asp Thr Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly 50 55 60

Ala Met Ser Asp Thr Arg Ile Asp Ala Ala Asp Arg Glu Ala Leu Gln 65 70 75 80

Ala Asn Leu Val Arg Ser His Ala Ala Gly Ala Gly Ser Glu Leu Asp 85 90 95

Thr Ala Ala Val Arg Ala Leu Leu Val Thr Arg Leu Asn Ala Leu Ala 100 105 110

Lys Gly Tyr Ser Gly Ile Arg Glu Arg Val Leu Asp Val Leu Val Gly

Leu Leu Asn Glu Gly Val His Pro Val Val Pro Ser Arg Gly Ser Leu 130 135 140

Gly Ala Ser Gly Asp Leu Ala Pro Leu Ala His Met Ser Arg Val Leu 145 150 155 160

Ile Gly Glu Gly Gln Ala Thr Asp Val Ala Gly Glu Arg Met Pro Ala 165 170 175

Ala Glu Ala Leu Ala Ala Ala Asp Leu Glu Pro Val Thr Leu Gln Ala 185 Lys Glu Gly Leu Ala Leu Ile Asn Gly Thr Gln Leu Thr Thr Gly Val Ala Ala Leu Ala Leu Val Asp Ala Glu Arg Val Leu Arg Ser Ala Asp 215 Thr Ala Gly Ala Leu Thr Thr Glu Val Thr Met Ser Thr Thr Ala Ser Cys Ala Pro Ala Ile His Glu Pro Val Arg Pro His Asp Gly Gln Ala 245 Val Ser Ala Arg His Ile Arg Asn Leu Thr Ala Gly Ser Glu Val Leu 260 265 Asp His His Arg Asp Cys Asp Gly Arg Val Gln Asp Ala Tyr Ser Ile 280 Arg Cys Leu Pro Gln Val His Gly Ala Val Arg Asp Ala Leu Asp His 290 295 300 Leu Arg Ala Ala Val Ala Thr Glu Leu Asn Ser Ala Thr Asp Asn Pro 310 315 Leu Val Phe Pro Ser Gly Thr Val Val Ser Gly Gly Asn Phe His Gly Glu Val Leu Ala Leu Arg Leu Gly Tyr Ala Ala Ser Ala Leu Ala Glu Leu Ala Ala Ile Ser Glu Arg Arg Thr Asp Arg Leu Leu Asn Pro Glu 360 Thr Gln Glu Pro Leu Glu Pro Phe Leu Ala Pro Asp Ser Gly Leu His 375 Ser Gly Leu Met Ile Pro Gln Tyr Thr Ala Ala Ser Leu Val Asn Asp 395 Leu Arg Ser Leu Gly Gln Pro Ala Thr Leu Asp Asn Ala Ser Val Ser 405 Gly Ala Gln Glu Asp His Val Ser Met Ser Ala Gly Ala Ala Tyr Asn 425 Phe Arg Glu Ala Val Glu Lys Ala Ala Thr Val Val Gly Val Glu Leu Leu Cys Gly Ala Gln Gly Arg Glu Phe Leu Asp Pro Leu Ala Leu Gly 455 Ala Gly Thr Ala Ala Ala Tyr Asp Leu Val Arg Ser Glu Val Ser Glu 475

Pro Ala Gly Asp Arg Ala Leu Ala Asp Asp Met Ala Ala Val Gly Asp 485 490 495

Leu Val Arg Ala Gly Leu Val Glu Asp Ala Val Ala Arg Ala Leu Asp 500 505 510

Ala

<210> 65

<211> 504

<212> PRT

<213> Streptomyces griseus

<400> 65

Val Val Val Gly Thr Ser Gly Thr Thr Ala Glu Asp Val Val Ala Val 1 5 10 15

Ala Arg His Gly Ala Arg Val Glu Leu Ser Ala Ala Ala Val Glu Ala 20 25 30

Leu Ala Ala Arg Leu Ile Val Asp Ala Leu Ala Ala Lys Pro Glu 35 40 45

Pro Val Tyr Gly Val Ser Thr Gly Phe Gly Ala Leu Ala Ser Arg His 50 55 60

Ile Gly Thr Glu Leu Arg Ala Gln Leu Gln Arg Asn Ile Val Arg Ser 65 70 75 80

His Ala Ala Gly Met Gly Pro Arg Val Glu Arg Glu Val Val Arg Ala $85\,$ 90 $95\,$

Leu Met Phe Leu Arg Leu Lys Thr Val Ala Ser Gly His Thr Gly Val 100 105 110

Arg Pro Glu Val Ala Gln Thr Met Ala Asp Val Leu Asn Ala Gly Ile 115 120 125

Thr Pro Val Val His Glu Tyr Gly Ser Leu Gly Cys Ser Gly Asp Leu 130 135 140

Ala Pro Leu Ser His Cys Ala Leu Thr Leu Met Gly Glu Gly Glu Ala 145 150 155 160

Glu Gly Pro Asp Gly Thr Val Arg Pro Ala Gly Glu Leu Leu Ala Ala 165 170 175

His Gly Ile Ala Pro Val Glu Leu Arg Glu Lys Glu Gly Leu Ala Leu 180 185 190

Leu Asn Gly Thr Asp Gly Met Leu Gly Met Leu Val Met Ala Leu Ala 195 200 205

Asp Leu Arg Asn Leu Tyr Thr Ser Ala Asp Ile Thr Ala Ala Leu Ser 210 215 220

Leu Glu Ala Leu Leu Gly Thr Asp Lys Val Leu Ala Pro Glu Leu His

235 Ala Ile Arg Pro His Pro Gly Gln Gly Val Ser Ala Asp Asn Met Ser Arg Val Leu Ala Gly Ser Gly Leu Thr Gly His His Gln Asp Asp Ala Pro Arg Val Gln Asp Ala Tyr Ser Val Arg Cys Ala Pro Gln Val Asn Gly Ala Gly Arg Asp Thr Leu Asp His Ala Ala Leu Val Ala Gly Arg 295 Glu Leu Ala Ser Ser Val Asp Asn Pro Val Val Leu Pro Asp Gly Arg 310 315 320 305 Val Glu Ser Asn Gly Asn Phe His Gly Ala Pro Val Ala Tyr Val Leu 325 330 Asp Phe Leu Ala Ile Val Ala Ala Asp Leu Gly Ser Ile Cys Glu Arg 340 345 Arg Thr Asp Arg Leu Leu Asp Lys Asn Arg Ser His Gly Leu Pro Pro 360 Phe Leu Ala Asp Asp Ala Gly Val Asp Ser Gly Leu Met Ile Ala Gln 375 Tyr Thr Gln Ala Ala Leu Val Ser Glu Met Lys Arg Leu Ala Val Pro 395 Ala Ser Ala Asp Ser Ile Pro Ser Ser Ala Met Gln Glu Asp His Val Ser Met Gly Trp Ser Ala Ala Arg Lys Leu Arg Thr Ala Val Asp Asn 425 Leu Ala Arg Ile Val Ala Val Glu Leu Tyr Ala Ala Thr Arg Ala Ile Glu Leu Arg Ala Ala Glu Gly Leu Thr Pro Ala Pro Ala Ser Glu Ala 455

Val Val Ala Ala Leu Arg Ala Ala Gly Ala Glu Gly Pro Gly Pro Asp

Arg Phe Leu Ala Pro Asp Leu Ala Ala Ala Asp Thr Phe Val Arg Glu

475

490

Gly Arg Leu Val Ala Ala Val Glu 500

470

<210> 66

<211> 502

<212> PRT

<213> Corynebacteriaceae sp.

Ala Arg His Glu Ala Arg Ile Ser Ile Ser Pro Gln Val Leu Glu Glu 20 25 30

Leu Ala Ser Val Arg Ala His Ile Asp Ala Leu Ala Ser Ala Asp Thr 35 40 45

Pro Val Tyr Gly Ile Ser Thr Gly Phe Gly Ala Leu Ala Thr Arg His 50 55 60

Ile Ala Pro Glu Asp Arg Ala Lys Leu Gln Arg Ser Leu Ile Arg Ser 65 70 75 80

His Ala Ala Gly Met Gly Glu Pro Val Glu Arg Glu Val Val Arg Ala 85 90 95

Leu Met Phe Leu Arg Ala Lys Thr Leu Ala Ser Gly Arg Thr Gly Val 100 105 110

Arg Pro Val Val Leu Glu Thr Met Val Gly Met Leu Asn Ala Gly Ile 115 120 125

Thr Pro Val Val Arg Glu Tyr Gly Ser Leu Gly Cys Ser Gly Asp Leu 130 135 140

Ala Pro Leu Ser His Cys Ala Leu Val Leu Met Gly Glu Gly Glu Ala 145 150 155 160

Thr Asp Ala His Gly Asp Ile Arg Pro Val Pro Glu Leu Phe Ala Glu 165 170 175

Ala Gly Leu Thr Pro Val Glu Leu Ala Glu Lys Glu Gly Leu Ala Leu 180 185 190

Val Asn Gly Thr Asp Gly Met Leu Gly Gln Leu Ile Met Ala Leu Ala 195 200 205

Asp Leu Asp Glu Leu Leu Asp Ile Ala Asp Ala Thr Ala Ala Met Ser 210 215 220

Val Glu Ala Gln Leu Gly Thr Asp Gln Val Phe Arg Ala Glu Leu His 225 230 235 240

Glu Pro Leu Arg Pro His Pro Gly Gln Gly Arg Ser Ala Gln Asn Met 245 250 255

Phe Ala Phe Leu Ala Asp Ser Pro Ile Val Ala Ser His Arg Glu Gly 260 265 270

Asp Gly Arg Val Gln Asp Ala Tyr Ser Leu Arg Cys Ser Pro Gln Val 275 280 285







Thr Gly Ala Ala Arg Asp Thr Ile Ala His Ala Arg Leu Val Ala Thr 295 Arg Glu Leu Ala Ala Ile Asp Asn Pro Val Val Leu Pro Ser Gly 315 Glu Val Thr Ser Asn Gly Asn Phe His Gly Ala Pro Val Ala Tyr Val Leu Asp Phe Leu Ala Ile Ala Val Ala Asp Leu Gly Ser Ile Ala Glu 340 Arg Arg Thr Asp Arg Met Leu Asp Pro Ala Arg Ser Arg Asp Leu Pro Ala Phe Leu Ala Asp Asp Pro Gly Val Asp Ser Gly Met Met Ile Ala 375 Gln Tyr Thr Gln Ala Gly Leu Val Ala Glu Asn Lys Arg Leu Ala Val 390 395 Pro Ala Ser Val Asp Ser Ile Pro Ser Ser Ala Met Gln Glu Asp His 410 415 Val Ser Leu Gly Trp His Ala Ala Arg Lys Leu Arg Thr Ser Val Ala Asn Leu Arg Arg Ile Leu Ala Val Glu Met Leu Ile Ala Gly Arg Ala Leu Asp Leu Arg Ala Pro Leu Lys Pro Gly Pro Ala Thr Gly Ala Val 455

Leu Glu Val Leu Arg Ser Lys Val Ala Gly Pro Gly Gln Asp Arg Phe

Leu Ser Ala Glu Leu Glu Ala Ala Tyr Asp Leu Leu Ala Asn Gly Ser 485 490 495

Val His Lys Ala Leu Glu 500

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